

SEQUENCE LISTING

<110> Albertsen, Hans
Anand, Rakesh
Carlson, Mary
Grodin, Joanna
Hedge, Philip John
Joslyn, Geoff
Kinzler, Kenneth
Markham, Alexander Fred
Nakamura, Yusuke
Thliveris, Andrew
Vogelstein, Bert
White, Raymond L.

<120> APC Antibodies

<130> 001107.78817

<140> US 09/442,489

<141> 1999-11-18

<150> US 08/452,654

<151> 1995-05-25

<150> US 08/289,548

<151> 1994-08-12

<150> US 07/741,940

<151> 1991-08-08

<160> 154

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 9606

<212> DNA

<213> Homo sapiens

<400> 1

ggactcggaa	atgaggtcca	agggtagcca	aggatggctg	cagcttcata	tgatcagttg	60
ttaaagcaag	ttgaggcact	gaagatggag	aactcaaata	ttcgacaaga	gctagaagat	120
aattccaatc	atcttacaata	actggaaaact	gagggatcta	atatgaagga	agtacttaaa	180
caactacaag	gaagtattga	agatgaagct	atggcttctt	ctggacagat	tgatttatta	240
gagcgtctta	aagagcttaa	cttagatagc	agtaatttcc	ctggagtaaa	actgagggtca	300
aaaatgtccc	tccgttctta	tggaaagccg	gaaggatctg	tatcaagccg	ttctggagag	360
tgcagtcctg	ttcctatggg	ttcatttcca	agaagagggt	ttgtaaatgg	aagcagagaa	420
agtactggat	atttagaaga	acttgagaaa	gagagggtcat	tgcttcttgc	tgatcttgac	480
aaagaagaaa	aggaaaaaga	ctggtattac	gctcaacttc	agaatctcac	taaaagaata	540
gatagtcttc	ctttaactga	aaatttttcc	ttacaaacag	atttgaccag	aaggcaattg	600
gaatatgaag	caaggcaaat	cagagttgag	atggaagaac	aactaggtac	ctgccaggat	660
atggaaaaac	gagcacagcg	aagaatagcc	agaattcagc	aaatcgaaaa	ggacataactt	720

cgtataccac	agctttttaca	gtcccaagca	acagaagcag	agaggtcatc	ttgaacaag	780
catgaaacog	gctcacatga	tgctgagcgg	cagaatgaag	gtcaaggagt	gggagaaatc	840
aacatggcaa	cttctggtaa	tggtcagggg	tcaactacac	gaatggacca	tgaaacagcc	900
agtgttttga	gttctagtag	cacacactct	gcacctcgaa	ggctgacaag	tcatctggga	960
accaaggtgg	aaatgggtga	ttcattgttg	tcaatgcttg	gtactcatga	taaggatgat	1020
atgtgcgcga	ctttgctagc	tatgtctagc	tcccaagaca	gctgtatatc	catgcgacag	1080
tctggatgtc	ttcctctcct	catccagctt	ttacatggca	atgacaaaga	ctctgtattg	1140
ttgggaaatt	cccggggcag	taaagaggct	cggggccagg	ccagtgcagc	actccacaac	1200
atcattcact	cadagcctga	tgacaagaga	ggcaggcgtg	aaatccgagt	ccttcatctt	1260
ttggaacaga	tacgcgctta	ctgtgaaacc	tggtgggagt	ggcagggaag	tcatgaacca	1320
ggcatggacc	aggacaaaaa	ttcaatgcca	gctcctgttg	aacatcagat	ctgtcctgct	1380
gtgtgtgttc	taatgaact	ttcatttgat	gaagagcata	gacatgcaat	gaatgaacta	1440
gggggactac	aggccattgc	agaattattg	caagtggact	gtgaaatgta	tgggcttact	1500
aatgaccact	acagtattac	actaagacga	tatgctggaa	tggctttgac	aaacttgact	1560
tttgagatg	tagccaacaa	ggctacgcga	tgctctatga	aaggctgcat	gagagcactt	1620
gtggcccaac	taaaatctga	aagtgaagac	ttacagcagg	ttattgcaag	tgttttgagg	1680
aattttctct	ggcgagcaga	tgtaaatagt	aaaaagacgt	tgcgagaagt	tggaaagtgtg	1740
aaagcattga	tggaaatgtgc	tttagaagtt	aaaaaggaat	caaccctcaa	aagcgtattg	1800
agtgccttat	ggaattttgtc	agcacattgc	actgagaata	aagctgatat	atgtgctgta	1860
gatggtgcac	ttgcattttt	gggtggcact	cttacttacc	ggagccagac	aaacacttta	1920
gccattattg	aaagtggagg	tgggatatta	cggaaatgtgt	ccagcttgat	agctacaaat	1980
gaggaccaca	ggcaaatcct	aagagagaac	aactgtctac	aaactttatt	acaacactta	2040
aaatctcata	gtttgacaat	agtcagtta	gcagtgtggaa	ctttgtggaa	tctctcagca	2100
agaaatccta	aagaccagga	agcattatgg	gacatggggg	cagttagcat	gctcaagaac	2160
ctcattcatt	caaagcacaa	aatgattgct	atgggaagt	ctgcagcttt	aaggaatctc	2220
atggcaata	ggcctgcgaa	gtacaaggat	gccaatatta	tgtctcctgg	ctcaagcttg	2280
ccatctcttc	atgttaggaa	acaaaaagcc	ctagaagcag	aattagatgc	tcagcactta	2340
tcagaaactt	ttgacaatat	agacaattta	agtcaccaagg	catctcatcg	tagtaagcag	2400
agacacaagc	aaagtctcta	tggtgattat	gtttttgaca	ccaatcgaca	tgatgataat	2460
aggtcagaca	attttaatac	tggaacacat	actgtccttt	caccatattt	gaatactaca	2520
gtgttaccca	gctcctcttc	atcaagagga	agcttagata	gttctcgttc	tgaaaaagat	2580
agaagtttgg	agagagaacg	cggaaattggt	ctaggcaact	accatccagc	aacagaaaaat	2640
ccaggaaactt	cttcaaagcg	aggtttgacg	atctccacca	ctgcagccca	gattgccaata	2700
gtcatggaag	aagtgtcagc	cattcatacc	tctcaggaag	acagaagttc	tgggtctacc	2760
actgaattac	attgtgtgac	agatgagaga	aatgcactta	gaagaagctc	tgctgccccat	2820
acacattcaa	acacttacaa	tttactaag	tcggaaaaat	caaataggac	atgttctatg	2880
ccttatgcc	aattagaata	caagagatct	tcaaatgata	gtttaaatag	tgtcagtagt	2940
aatgatggtt	atggtaaaag	agggtcaaatg	aaaccctcga	ttgaatecta	ttctgaagat	3000
gatgaaagta	agttttgacg	ttatggtcaa	taccagcccg	actagccca	tataatcat	3060
agtgcaaate	atatggatga	taatgatgga	gaactagata	caaccaataaa	ttatagtctt	3120
aaatattcag	atgagcagtt	gaactctgga	aggcaaagtc	cttcacagaa	tgaaagatgg	3180
gcaagaccga	aacacataat	agaagatgaa	ataaaacaaa	gtgagcaaa	acaatcaagg	3240
aatcaaagta	caacttatcc	tgttttatact	gagagcactg	atgataaaca	cctcaagttc	3300
caaccacatt	ttggacagca	ggaatgtgtt	tctccataca	ggtcacgggg	agccaatggt	3360
tcagaaacaa	atcgagtggg	ttctaatacat	ggaattaatc	aaaatgtaag	ccagtctttg	3420
tgtcaagaag	atgactatga	agatgataag	cctaccaatt	atagtgaacg	ttactctgaa	3480
gaagaacacg	atgaagaaga	agagagacca	acaaattata	gcataaaata	taatgaagag	3540
aaacgtcatg	tgatcagcc	tattgattat	agtttaaaat	atgccacaga	tattccttca	3600
tcacagaaac	agtcattttc	attctcaaag	agttcatctg	gacaaagcag	taaaaccgaa	3660
catatgtctt	caagcagtg	gaatacgtcc	acaccttcat	ctaatagccaa	gaggcagaat	3720
cagctccatc	caagttctgc	acagagtaga	agtggtcagc	ctcaaaaggc	tgocacttgc	3780
aaagtttctt	ctattaacca	agaaacaata	cagacttatt	gtgtagaaga	tactccaata	3840
tgtttttcaa	gatgtagtgc	attatcatct	ttgtcatcag	ctgaagatga	aataggatgt	3900
aatcagacga	cacaggaagc	agattctgct	aataccctgc	aaatagcaga	aataaaagga	3960
aagattggaa	ctaggtcagc	tgaagatcct	gtgagcgaag	ttccagcagt	gtcacagcac	4020

cctagaacca	aatccagcag	atgagcaggt	tctagtttat	cttcagaatc	agccaggcac	4080
aaagctgttg	aatctccttc	aggagcgaaa	tctccctcca	aaagtgggtgc	tcagacaccc	4140
aaaagtcac	ctgaacacta	tggttcaggag	acccactca	tgtttagcag	atgtacttct	4200
gtcagttcac	ttgatagttt	tgagagtcgt	tgcattgcca	gctccgttca	gagtgaacca	4260
tgcagtggaa	tggttaagtgg	cattataagc	cccagtgatc	ttccagatag	ccctggacaa	4320
accatgccac	caagcagaag	taaaacacct	ccaccacctc	ctcaaacagc	tcaaaccaag	4380
cgagaagtac	ctaaaaataa	agcacctact	gctgaaaaga	gagagagtgg	acctaagcaa	4440
gctgcagtaa	atgctgcagt	tcagaggggtc	caggttcttc	cagatgctga	tactttatta	4500
cattttgcca	cagaaagtas	tccagatgga	ttttcttggt	catccagcct	gagtgtctctg	4560
agcctcgatg	agccattttat	acagaaaagat	gtggaattaa	gaataatgcc	tccagttcag	4620
gaaaatgaca	atgggaatga	aacagaatca	gagcagccta	aagaatcaaa	tgaaaaccaa	4680
gagaaagagg	cagaaaaaac	tattgattct	gaaaaggacc	tattagatga	ttcagatgat	4740
gatgatattg	aaatactaga	agaatgtatt	atctctgcca	tgccaacaaa	gtcatcacgt	4800
aaaggcaaaa	agccagccca	gactgcttca	aaattacctc	cacctgtggc	aaggaaacca	4860
agtcagctgc	ctgtgtacaa	acttctacca	tcacaaaaca	ggttgcaacc	ccaaaagcat	4920
gtagtgttta	caccggggga	tgatatgcca	cgggtgtatt	gtgttgaaag	gacacctata	4980
aaacttttcca	cagctacatc	tctaagtgtg	ctaacaatcg	aatccctctc	aaatgagtta	5040
gctgctggag	aaggagttag	aggaggagca	cagtcagggtg	aatgtgaaaa	acgagatacc	5100
attcctacag	aaggcagaag	tacagatgag	gctcaaggag	gaaaaacctc	atctgtaacc	5160
atacctgaat	tgatgacaa	taaagcagag	gaaggtgata	ttcttgacaga	atgcattaat	5220
tctgctatgc	ccaaagggaa	aagtcacaag	cctttccgtg	tgaaaaagat	aatggaccag	5280
gtccagcaag	catctgcgtc	gtcttctgca	cccaacaaaa	atcagttaga	tggttaagaa	5340
aagaaaccaa	cttcaccagt	aaaacctata	ccacaaaata	ctgaatatag	gacacgtgtg	5400
agaaaaaatg	cagactcaaa	aaataattta	aatgctgaga	gagttttctc	agacaacaaa	5460
gattcaaaga	aacagaatct	gaaaaataat	tccaaggact	tcaatgataa	gctcccaaat	5520
aatgaagata	gagtcagagg	aagttttgct	tttgattcac	ctcatcatta	cacgcctatt	5580
gaaggaactc	cttactgttt	ttcacgaaat	gattctttga	gttctctaga	ttttgatgat	5640
gatgatgttg	acctttccag	ggaaaaggct	gaattaagaa	aggcaaaaga	aaataaggaa	5700
tcagaggcta	aagttaccag	ccacacagaa	ctaacctcca	accaacaatc	agctaataag	5760
acacaagcta	ttgcaaaagca	gccaataaat	cgaggtcagc	ctaaacccat	acttcagaaa	5820
caatccactt	ttccccagtc	atccaaagac	ataccagaca	gaggggcagc	aactgatgaa	5880
aagttacaga	atatttgctat	tgaaaaatact	ccagtttgct	tttctcataa	ttcctctctg	5940
agttctctca	gtgacattga	ccaagaaaac	aacaataaag	aaaatgaacc	tatcaaagag	6000
actgagcccc	ctgactcaca	gggagaacca	agtaaacctc	aagcatcagg	ctatgtctct	6060
aaatcatttc	atgttgaaaga	taccccagtt	tgttttctca	gaacacagtc	tctcagttct	6120
cttagtattg	actctgaaga	tgacctgttg	caggaatgta	taagctccgc	aatgccaaaa	6180
aagaaaaagc	cttcaagact	caagggtgat	aatgaaaaac	atagtcccag	aaatatgggt	6240
ggcatattag	gtgaagatct	gacacttgat	ttgaaagata	tacagagacc	agattcagaa	6300
catgggtctat	cccttgattc	agaaaatttt	gattggaaaag	ctattcagga	aggtgcaaat	6360
tccatagtaa	gtagttttaca	tcaagctgct	gctgctgcat	gtttatctag	acaagcttcg	6420
tctgattcag	attccatcct	ttccctgaaa	tcaggaatct	ctctgggac	accatttcat	6480
cttacacctg	atcaagaaga	aaaacccttt	acaagtaata	aaggcccaag	aattctaaaa	6540
ccaggggaga	aaagtacatt	ggaaactaaa	aagatagaat	ctgaaagtaa	aggaatcaaa	6600
ggaggaaaaaa	aagtttataa	aagtttgatt	actggaaaag	ttcgatctaa	ttcagaaatt	6660
tcaggccaaa	tgaaacagcc	ccttcaagca	aacatgcctt	caatctctcg	aggcaggaca	6720
atgattcata	ttccaggagt	tcgaaatagc	tcttcaagta	caagtcctgt	ttctaaaaaa	6780
ggccaccccc	ttaagactcc	agcctccaaa	agccttagtg	aagggtcaaac	agccaccact	6840
tctcctagag	gagccaagcc	atctgtgaaa	tcagaattaa	gccctgttgc	cagccagaca	6900
tcccaaatag	gtgggtcaag	taaagcacct	tctagatcag	gatctagaga	ttcgacccct	6960
tcaagacctg	cccagcaacc	attaagtaga	cctatacagt	ctcctggccg	aaactcaatt	7020
tcccctggta	gaaatggaat	aagtcctcct	aacaaattat	ctcaacttcc	aaggacatca	7080
tcccctagta	ctgcttcaac	taagtcctca	ggttctggaa	aaatgtcata	tacatctcca	7140
ggtagacaga	tgagccaaca	gaaccttacc	aaacaaacag	gtttatccaa	gaatgccagt	7200
agtatttcaa	gaagtgaagtc	tgccctccaaa	ggactaaatc	agatgaataa	tggtaatgga	7260
gccaataaaa	aggtagaact	ttctagaatg	tcttcaacta	aatcaagtg	aagtgaatct	7320

gatagatcag aaagacctgt attagtagcgc cagtcactt tcatcaaaga agctccaagc 7380
 ccaaccttaa gaagaaaatt ggaggaatct gcttcatttg aatctctttc tccatcatct 7440
 agaccagctt ctccactag gtcccaggca caaactccag ttttaagtc tcccttccct 7500
 gatattgtctc tatccacaca ttctgtctgt caggctggtg gatggcgaaa actcccacct 7560
 aatctcagtc ccaatataga gtataatgat ggaagaccag caaagcgcca tgatattgca 7620
 cgggtctcatt ctgaagtcct ttctagactt ccaatcaata ggtcaggaac ctggaaacgt 7680
 gagcacagca aacattcatc atcccttccct cgagtaagca cttggagaag aactggaagt 7740
 tcatcttcaa ttctttctgc ttcatcagaa tccagtgaag aagcaaaaag tgaggatgaa 7800
 aaacatgtga actctatttc aggaacccaa caaagtaaaag aaaaccaagt atccgcaaaa 7860
 ggaacatgga gaaaaataaa agaaaatgaa ttttctcca caaatagtac ttctcagacc 7920
 gtttccctcag gtgctacaaa tgggtgctgaa tcaaagactc taatttatca aatggcacct 7980
 gctgtttcta aaacagagga tggttggtg agaattgagg actgtcccat taacaatcct 8040
 agatctggaa gatctccac aggttaatact ccccggtga ttgacagtgt ttcagaaaag 8100
 gcaaatccaa acattaaaga ttcaaaagat aatcaggcaa aacaaaatgt gggtaatggc 8160
 agtggtccca tgcgtaccgt gggtttgaa aatgcctga cctcctttat tcaggtggat 8220
 gccctgacc aaaaaggaac tgagataaaa ccaggacaaa ataactctgt ccctgtatca 8280
 gagactaat aaagtcctat agtggaaagt accccattca gttctagcag ctcaagcaaa 8340
 cacagttcac ctagtgggac tgggtgctgc agagtgactc cttttaatta caaccaagc 8400
 cctaggaaaa gcagcgcaga tagcacttca gctcggccat ctcagatccc aactccagt 8460
 aataacaaca caaagaagcg agattccaaa actgacagca cagaatccag tggaaaccaa 8520
 agtcctaagc gccattctgg gtcttacctt gtgacatctg tttaaaagag aggaagaatg 8580
 aaactaagaa aattctatgt taattacaac tgctatatag acattttgtt tcaaatgaaa 8640
 ctttaaaaaga ctgaaaaatt ttgtatatag gtttgattct tgtagaggg tttttgttct 8700
 ggaagccata tttgatagta tacttttct tcaactggtct tttttggga ggcactcttg 8760
 atggttagga aaaaatagaa agccaagtat gttgtacag tatgttttac atgtatttaa 8820
 agtagcatcc catcccaact tccttaatta ttgcttgctt aaaataatga acactacaga 8880
 taggaaatat gatatttgc ttttatcaat catttctaga ttataaactg actaaactta 8940
 catcagggga aaattgggtat ttatgcaaaa aaaaaatgtt tttgtccttg tgagtcctc 9000
 taacatcata attaatcatg tggctgtgaa attcacagta atatggttcc cgatgaacaa 9060
 gtttaccag cctgcttgc ttactgcagt aatgaaactg atggttcaat ttcagaagta 9120
 atgattaaca gttatgtgt cacatgatgt gcataagat agctacagt taataattta 9180
 cactattttg tgctccaac aaaacaaaaa tctgtgtaac tgtaaaacat tgaatgaaac 9240
 tattttacct gaactagatt ttatctgaaa gtaggtagaa tttttgctat gctgtaattt 9300
 gttgtatatt ctggtatttg aggtgagatg gctgtctttt attaatgaga catgaattgt 9360
 gtctcaacag aaactaaatg aacatttcag aataaattat tgctgtatgt aaactgttac 9420
 tgaaattggt atttgtttga aggttttgtt tcacatttgt attaatat tgtttaaaat 9480
 gcctctttta aaagcttata taaatttttt cttcagcttc tatgcattaa gagtaaaatt 9540
 cctcttactg taataaaaaac attgaagaag actgttgcca cttaaccatt ccatgcgttg 9600
 gcactt

<210> 2
 <211> 2843
 <212> PRT
 <213> Homo sapiens

<400> 2
 Met Ala Ala Ala Ser Tyr Asp Gln Leu Leu Lys Gln Val Glu Ala Leu
 1 5 10 15
 Lys Met Glu Asn Ser Asn Leu Arg Gln Glu Leu Glu Asp Asn Ser Asn
 20 25 30
 His Leu Thr Lys Leu Glu Thr Glu Ala Ser Asn Met Lys Glu Val Leu
 35 40 45
 Lys Gln Leu Gln Gly Ser Ile Glu Asp Glu Ala Met Ala Ser Ser Gly

50 55 60
 Gln Ile Asp Leu Leu Glu Arg Leu Lys Glu Leu Asn Leu Asp Ser Ser
 65 70 75 80
 Asn Phe Pro Gly Val Lys Leu Arg Ser Lys Met Ser Leu Arg Ser Tyr
 85 90 95
 Gly Ser Arg Glu Gly Ser Val Ser Ser Arg Ser Gly Glu Cys Ser Pro
 100 105 110
 Val Pro Met Gly Ser Phe Pro Arg Gly Phe Val Asn Gly Ser Arg
 115 120 125
 Glu Ser Thr Gly Tyr Leu Glu Glu Leu Glu Lys Glu Arg Ser Leu Leu
 130 135 140
 Leu Ala Asp Leu Asp Lys Glu Glu Lys Glu Lys Asp Trp Tyr Tyr Ala
 145 150 155 160
 Gln Leu Gln Asn Leu Thr Lys Arg Ile Asp Ser Leu Pro Leu Thr Glu
 165 170 175
 Asn Phe Ser Leu Gln Thr Asp Leu Thr Arg Arg Gln Leu Glu Tyr Glu
 180 185 190
 Ala Arg Gln Ile Arg Val Ala Met Glu Glu Gln Leu Gly Thr Cys Gln
 195 200 205
 Asp Met Glu Lys Arg Ala Gln Arg Arg Ile Ala Arg Ile Gln Gln Ile
 210 215 220
 Glu Lys Asp Ile Leu Arg Ile Arg Gln Leu Leu Gln Ser Gln Ala Thr
 225 230 235 240
 Glu Ala Glu Arg Ser Ser Gln Asn Lys His Glu Thr Gly Ser His Asp
 245 250 255
 Ala Glu Arg Gln Asn Glu Gly Gln Gly Val Gly Glu Ile Asn Met Ala
 260 265 270
 Thr Ser Gly Asn Gly Gln Gly Ser Thr Thr Arg Met Asp His Glu Thr
 275 280 285
 Ala Ser Val Leu Ser Ser Ser Ser Thr His Ser Ala Pro Arg Arg Leu
 290 295 300
 Thr Ser His Leu Gly Thr Lys Val Glu Met Val Tyr Ser Leu Leu Ser
 305 310 315 320
 Met Leu Gly Thr His Asp Lys Asp Asp Met Ser Arg Thr Leu Leu Ala
 325 330 335
 Met Ser Ser Ser Gln Asp Ser Cys Ile Ser Met Arg Gln Ser Gly Cys
 340 345 350
 Leu Pro Leu Leu Ile Gln Leu Leu His Gly Asn Asp Lys Asp Ser Val
 355 360 365
 Leu Leu Gly Asn Ser Arg Gly Ser Lys Glu Ala Arg Ala Arg Ala Ser
 370 375 380
 Ala Ala Leu His Asn Ile Ile His Ser Gln Pro Asp Asp Lys Arg Gly
 385 390 395 400
 Arg Arg Glu Ile Arg Val Leu His Leu Leu Glu Gln Ile Arg Ala Tyr
 405 410 415
 Cys Glu Thr Cys Trp Glu Trp Gln Glu Ala His Glu Pro Gly Met Asp
 420 425 430
 Gln Asp Lys Asn Pro Met Pro Ala Pro Val Glu His Gln Ile Cys Pro
 435 440 445
 Ala Val Cys Val Leu Met Lys Leu Ser Phe Asp Glu Glu His Arg His
 450 455 460
 Ala Met Asn Glu Leu Gly Gly Leu Gln Ala Ile Ala Glu Leu Leu Gln
 465 470 475 480
 Val Asp Cys Glu Met Tyr Gly Leu Thr Asn Asp His Tyr Ser Ile Thr
 485 490 495

Leu Arg Arg Tyr Ala Gly Met Ala Leu Thr Asn Leu Thr Phe Gly Asp
 500 505 510
 Val Ala Asn Lys Ala Thr Leu Cys Ser Met Lys Gly Cys Met Arg Ala
 515 520 525
 Leu Val Ala Gln Leu Lys Ser Glu Ser Glu Asp Leu Gln Gln Val Ile
 530 535 540
 Ala Ser Val Leu Arg Asn Leu Ser Trp Arg Ala Asp Val Asn Ser Lys
 545 550 555 560
 Lys Thr Leu Arg Glu Val Gly Ser Val Lys Ala Leu Met Glu Cys Ala
 565 570 575
 Leu Glu Val Lys Lys Glu Ser Thr Leu Lys Ser Val Leu Ser Ala Leu
 580 585 590
 Trp Asn Leu Ser Ala His Cys Thr Glu Asn Lys Ala Asp Ile Cys Ala
 595 600 605
 Val Asp Gly Ala Leu Ala Phe Leu Val Gly Thr Leu Thr Tyr Arg Ser
 610 615 620
 Gln Thr Asn Thr Leu Ala Ile Ile Glu Ser Gly Gly Gly Ile Leu Arg
 625 630 635 640
 Asn Val Ser Ser Leu Ile Ala Thr Asn Glu Asp His Arg Gln Ile Leu
 645 650 655
 Arg Glu Asn Asn Cys Leu Gln Thr Leu Leu Gln His Leu Lys Ser His
 660 665 670
 Ser Leu Thr Ile Val Ser Asn Ala Cys Gly Thr Leu Trp Asn Leu Ser
 675 680 685
 Ala Arg Asn Pro Lys Asp Gln Glu Ala Leu Trp Asp Met Gly Ala Val
 690 695 700
 Ser Met Leu Lys Asn Leu Ile His Ser Lys His Lys Met Ile Ala Met
 705 710 715 720
 Gly Ser Ala Ala Ala Leu Arg Asn Leu Met Ala Asn Arg Pro Ala Lys
 725 730 735
 Tyr Lys Asp Ala Asn Ile Met Ser Pro Gly Ser Ser Leu Pro Ser Leu
 740 745 750
 His Val Arg Lys Gln Lys Ala Leu Glu Ala Glu Leu Asp Ala Gln His
 755 760 765
 Leu Ser Glu Thr Phe Asp Asn Ile Asp Asn Leu Ser Pro Lys Ala Ser
 770 775 780
 His Arg Ser Lys Gln Arg His Lys Gln Ser Leu Tyr Gly Asp Tyr Val
 785 790 795 800
 Phe Asp Thr Asn Arg His Asp Asp Asn Arg Ser Asp Asn Phe Asn Thr
 805 810 815
 Gly Asn Met Thr Val Leu Ser Pro Tyr Leu Asn Thr Thr Val Leu Pro
 820 825 830
 Ser Ser Ser Ser Ser Arg Gly Ser Leu Asp Ser Ser Arg Ser Glu Lys
 835 840 845
 Asp Arg Ser Leu Glu Arg Glu Arg Gly Ile Gly Leu Gly Asn Tyr His
 850 855 860
 Pro Ala Thr Glu Asn Pro Gly Thr Ser Ser Lys Arg Gly Leu Gln Ile
 865 870 875 880
 Ser Thr Thr Ala Ala Gln Ile Ala Lys Val Met Glu Glu Val Ser Ala
 885 890 895
 Ile His Thr Ser Gln Glu Asp Arg Ser Ser Gly Ser Thr Thr Glu Leu
 900 905 910
 His Cys Val Thr Asp Glu Arg Asn Ala Leu Arg Arg Ser Ser Ala Ala
 915 920 925
 His Thr His Ser Asn Thr Tyr Asn Phe Thr Lys Ser Glu Asn Ser Asn

930
 Arg Thr Cys Ser Met Pro Tyr Ala Lys Leu Glu Tyr Lys Arg Ser Ser
 945 950 955 960
 Asn Asp Ser Leu Asn Ser Val Ser Ser Asn Asp Gly Tyr Gly Lys Arg
 965 970 975
 Gly Gln Met Lys Pro Ser Ile Glu Ser Tyr Ser Glu Asp Asp Glu Ser
 980 985 990
 Lys Phe Cys Ser Tyr Gly Gln Tyr Pro Ala Asp Leu Ala His Lys Ile
 995 1000 1005
 His Ser Ala Asn His Met Asp Asp Asn Asp Gly Glu Leu Asp Thr Pro
 1010 1015 1020
 Ile Asn Tyr Ser Leu Lys Tyr Ser Asp Glu Gln Leu Asn Ser Gly Arg
 1025 1030 1035 1040
 Gln Ser Pro Ser Gln Asn Glu Arg Trp Ala Arg Pro Lys His Ile Ile
 1045 1050 1055
 Glu Asp Glu Ile Lys Gln Ser Glu Gln Arg Gln Ser Arg Asn Gln Ser
 1060 1065 1070
 Thr Thr Tyr Pro Val Tyr Thr Glu Ser Thr Asp Asp Lys His Leu Lys
 1075 1080 1085
 Phe Gln Pro His Phe Gly Gln Gln Glu Cys Val Ser Pro Tyr Arg Ser
 1090 1095 1100
 Arg Gly Ala Asn Gly Ser Glu Thr Asn Arg Val Gly Ser Asn His Gly
 1105 1110 1115 1120
 Ile Asn Gln Asn Val Ser Gln Ser Leu Cys Gln Glu Asp Asp Tyr Glu
 1125 1130 1135
 Asp Asp Lys Pro Thr Asn Tyr Ser Glu Arg Tyr Ser Glu Glu Glu Gln
 1140 1145 1150
 His Glu Glu Glu Glu Arg Pro Thr Asn Tyr Ser Ile Lys Tyr Asn Glu
 1155 1160 1165
 Glu Lys Arg His Val Asp Gln Pro Ile Asp Tyr Ser Leu Lys Tyr Ala
 1170 1175 1180
 Thr Asp Ile Pro Ser Ser Gln Lys Gln Ser Phe Ser Phe Ser Lys Ser
 1185 1190 1195 1200
 Ser Ser Gly Gln Ser Ser Lys Thr Glu His Met Ser Ser Ser Ser Glu
 1205 1210 1215
 Asn Thr Ser Thr Pro Ser Ser Asn Ala Lys Arg Gln Asn Gln Leu His
 1220 1225 1230
 Pro Ser Ser Ala Gln Ser Arg Ser Gly Gln Pro Gln Lys Ala Ala Thr
 1235 1240 1245
 Cys Lys Val Ser Ser Ile Asn Gln Glu Thr Ile Gln Thr Tyr Cys Val
 1250 1255 1260
 Glu Asp Thr Pro Ile Cys Phe Ser Arg Cys Ser Ser Leu Ser Ser Leu
 1265 1270 1275 1280
 Ser Ser Ala Glu Asp Glu Ile Gly Cys Asn Gln Thr Thr Gln Glu Ala
 1285 1290 1295
 Asp Ser Ala Asn Thr Leu Gln Ile Ala Glu Ile Lys Gly Lys Ile Gly
 1300 1305 1310
 Thr Arg Ser Ala Glu Asp Pro Val Ser Glu Val Pro Ala Val Ser Gln
 1315 1320 1325
 His Pro Arg Thr Lys Ser Ser Arg Leu Gln Gly Ser Ser Leu Ser Ser
 1330 1335 1340
 Glu Ser Ala Arg His Lys Ala Val Glu Phe Pro Ser Gly Ala Lys Ser
 1345 1350 1355 1360
 Pro Ser Lys Ser Gly Ala Gln Thr Pro Lys Ser Pro Pro Glu His Tyr
 1365 1370 1375

Val Gln Glu Thr Pro Leu Met Phe Ser Arg Cys Thr Ser Val Ser Ser
 1380 1385 1390
 Leu Asp Ser Phe Glu Ser Arg Ser Ile Ala Ser Ser Val Gln Ser Glu
 1395 1400 1405
 Pro Cys Ser Gly Met Val Ser Gly Ile Ile Ser Pro Ser Asp Leu Pro
 1410 1415 1420
 Asp Ser Pro Gly Gln Thr Met Pro Pro Ser Arg Ser Lys Thr Pro Pro
 1425 1430 1435 1440
 Pro Pro Pro Gln Thr Ala Gln Thr Lys Arg Glu Val Pro Lys Asn Lys
 1445 1450 1455
 Ala Pro Thr Ala Glu Lys Arg Glu Ser Gly Pro Lys Gln Ala Ala Val
 1460 1465 1470
 Asn Ala Ala Val Gln Arg Val Gln Val Leu Pro Asp Ala Asp Thr Leu
 1475 1480 1485
 Leu His Phe Ala Thr Glu Ser Thr Pro Asp Gly Phe Ser Cys Ser Ser
 1490 1495 1500
 Ser Leu Ser Ala Leu Ser Leu Asp Glu Pro Phe Ile Gln Lys Asp Val
 1505 1510 1515 1520
 Glu Leu Arg Ile Met Pro Pro Val Gln Glu Asn Asp Asn Gly Asn Glu
 1525 1530 1535
 Thr Glu Ser Glu Gln Pro Lys Glu Ser Asn Glu Asn Gln Glu Lys Glu
 1540 1545 1550
 Ala Glu Lys Thr Ile Asp Ser Glu Lys Asp Leu Leu Asp Asp Ser Asp
 1555 1560 1565
 Asp Asp Asp Ile Glu Ile Leu Glu Glu Cys Ile Ile Ser Ala Met Pro
 1570 1575 1580
 Thr Lys Ser Ser Arg Lys Gly Lys Lys Pro Ala Gln Thr Ala Ser Lys
 1585 1590 1595 1600
 Leu Pro Pro Pro Val Ala Arg Lys Pro Ser Gln Leu Pro Val Tyr Lys
 1605 1610 1615
 Leu Leu Pro Ser Gln Asn Arg Leu Gln Pro Gln Lys His Val Ser Phe
 1620 1625 1630
 Thr Pro Gly Asp Asp Met Pro Arg Val Tyr Cys Val Glu Gly Thr Pro
 1635 1640 1645
 Ile Asn Phe Ser Thr Ala Thr Ser Leu Ser Asp Leu Thr Ile Glu Ser
 1650 1655 1660
 Pro Pro Asn Glu Leu Ala Ala Gly Glu Gly Val Arg Gly Gly Ala Gln
 1665 1670 1675 1680
 Ser Gly Glu Phe Glu Lys Arg Asp Thr Ile Pro Thr Glu Gly Arg Ser
 1685 1690 1695
 Thr Asp Glu Ala Gln Gly Gly Lys Thr Ser Ser Val Thr Ile Pro Glu
 1700 1705 1710
 Leu Asp Asp Asn Lys Ala Glu Glu Gly Asp Ile Leu Ala Glu Cys Ile
 1715 1720 1725
 Asn Ser Ala Met Pro Lys Gly Lys Ser His Lys Pro Phe Arg Val Lys
 1730 1735 1740
 Lys Ile Met Asp Gln Val Gln Gln Ala Ser Ala Ser Ser Ser Ala Pro
 1745 1750 1755 1760
 Asn Lys Asn Gln Leu Asp Gly Lys Lys Lys Lys Pro Thr Ser Pro Val
 1765 1770 1775
 Lys Pro Ile Pro Gln Asn Thr Glu Tyr Arg Thr Arg Val Arg Lys Asn
 1780 1785 1790
 Ala Asp Ser Lys Asn Asn Leu Asn Ala Glu Arg Val Phe Ser Asp Asn
 1795 1800 1805
 Lys Asp Ser Lys Lys Gln Asn Leu Lys Asn Asn Ser Lys Asp Phe Asn

1810 1815 1820
 Asp Lys Leu Pro Asn Asn Glu Asp Arg Val Arg Gly Ser Phe Ala Phe
 1825 1830 1835 1840
 Asp Ser Pro His His Tyr Thr Pro Ile Glu Gly Thr Pro Tyr Cys Phe
 1845 1850 1855
 Ser Arg Asn Asp Ser Leu Ser Ser Leu Asp Phe Asp Asp Asp Val
 1860 1865 1870
 Asp Leu Ser Arg Glu Lys Ala Glu Leu Arg Lys Ala Lys Glu Asn Lys
 1875 1880 1885
 Glu Ser Glu Ala Lys Val Thr Ser His Thr Glu Leu Thr Ser Asn Gln
 1890 1895 1900
 Gln Ser Ala Asn Lys Thr Gln Ala Ile Ala Lys Gln Pro Ile Asn Arg
 1905 1910 1915 1920
 Gly Gln Pro Lys Pro Ile Leu Gln Lys Gln Ser Thr Phe Pro Gln Ser
 1925 1930 1935
 Ser Lys Asp Ile Pro Asp Arg Gly Ala Ala Thr Asp Glu Lys Leu Gln
 1940 1945 1950
 Asn Phe Ala Ile Glu Asn Thr Pro Val Cys Phe Ser His Asn Ser Ser
 1955 1960 1965
 Leu Ser Ser Leu Ser Asp Ile Asp Gln Glu Asn Asn Asn Lys Glu Asn
 1970 1975 1980
 Glu Pro Ile Lys Glu Thr Glu Pro Pro Asp Ser Gln Gly Glu Pro Ser
 1985 1990 1995 2000
 Lys Pro Gln Ala Ser Gly Tyr Ala Pro Lys Ser Phe His Val Glu Asp
 2005 2010 2015
 Thr Pro Val Cys Phe Ser Arg Asn Ser Ser Leu Ser Ser Leu Ser Ile
 2020 2025 2030
 Asp Ser Glu Asp Asp Leu Leu Gln Glu Cys Ile Ser Ser Ala Met Pro
 2035 2040 2045
 Lys Lys Lys Lys Pro Ser Arg Leu Lys Gly Asp Asn Glu Lys His Ser
 2050 2055 2060
 Pro Arg Asn Met Gly Gly Ile Leu Gly Glu Asp Leu Thr Leu Asp Leu
 2065 2070 2075 2080
 Lys Asp Ile Gln Arg Pro Asp Ser Glu His Gly Leu Ser Pro Asp Ser
 2085 2090 2095
 Glu Asn Phe Asp Trp Lys Ala Ile Gln Glu Gly Ala Asn Ser Ile Val
 2100 2105 2110
 Ser Ser Leu His Gln Ala Ala Ala Ala Ala Cys Leu Ser Arg Gln Ala
 2115 2120 2125
 Ser Ser Asp Ser Asp Ser Ile Leu Ser Leu Lys Ser Gly Ile Ser Leu
 2130 2135 2140
 Gly Ser Pro Phe His Leu Thr Pro Asp Gln Glu Glu Lys Pro Phe Thr
 2145 2150 2155 2160
 Ser Asn Lys Gly Pro Arg Ile Leu Lys Pro Gly Glu Lys Ser Thr Leu
 2165 2170 2175
 Glu Thr Lys Lys Ile Glu Ser Glu Ser Lys Gly Ile Lys Gly Gly Lys
 2180 2185 2190
 Lys Val Tyr Lys Ser Leu Ile Thr Gly Lys Val Arg Ser Asn Ser Glu
 2195 2200 2205
 Ile Ser Gly Gln Met Lys Gln Pro Leu Gln Ala Asn Met Pro Ser Ile
 2210 2215 2220
 Ser Arg Gly Arg Thr Met Ile His Ile Pro Gly Val Arg Asn Ser Ser
 2225 2230 2235 2240
 Ser Ser Thr Ser Pro Val Ser Lys Lys Gly Pro Pro Leu Lys Thr Pro
 2245 2250 2255

Ala Ser Lys Ser Pro Ser Glu Gly Gln Thr Ala Thr Thr Ser Pro Arg
 2260 2265 2270
 Gly Ala Lys Pro Ser Val Lys Ser Glu Leu Ser Pro Val Ala Arg Gln
 2275 2280 2285
 Thr Ser Gln Ile Gly Gly Ser Ser Lys Ala Pro Ser Arg Ser Gly Ser
 2290 2295 2300
 Arg Asp Ser Thr Pro Ser Arg Pro Ala Gln Gln Pro Leu Ser Arg Pro
 2305 2310 2315 2320
 Ile Gln Ser Pro Gly Arg Asn Ser Ile Ser Pro Gly Arg Asn Gly Ile
 2325 2330 2335
 Ser Pro Pro Asn Lys Leu Ser Gln Leu Pro Arg Thr Ser Ser Pro Ser
 2340 2345 2350
 Thr Ala Ser Thr Lys Ser Ser Gly Ser Gly Lys Met Ser Tyr Thr Ser
 2355 2360 2365
 Pro Gly Arg Gln Met Ser Gln Gln Asn Leu Thr Lys Gln Thr Gly Leu
 2370 2375 2380
 Ser Lys Asn Ala Ser Ser Ile Pro Arg Ser Glu Ser Ala Ser Lys Gly
 2385 2390 2395 2400
 Leu Asn Gln Met Asn Asn Gly Asn Gly Ala Asn Lys Lys Val Glu Leu
 2405 2410 2415
 Ser Arg Met Ser Ser Thr Lys Ser Ser Gly Ser Glu Ser Asp Arg Ser
 2420 2425 2430
 Glu Arg Pro Val Leu Val Arg Gln Ser Thr Phe Ile Lys Glu Ala Pro
 2435 2440 2445
 Ser Pro Thr Leu Arg Arg Lys Leu Glu Glu Ser Ala Ser Phe Glu Ser
 2450 2455 2460
 Leu Ser Pro Ser Ser Arg Pro Ala Ser Pro Thr Arg Ser Gln Ala Gln
 2465 2470 2475 2480
 Thr Pro Val Leu Ser Pro Ser Leu Pro Asp Met Ser Leu Ser Thr His
 2485 2490 2495
 Ser Ser Val Gln Ala Gly Gly Trp Arg Lys Leu Pro Pro Asn Leu Ser
 2500 2505 2510
 Pro Thr Ile Glu Tyr Asn Asp Gly Arg Pro Ala Lys Arg His Asp Ile
 2515 2520 2525
 Ala Arg Ser His Ser Glu Ser Pro Ser Arg Leu Pro Ile Asn Arg Ser
 2530 2535 2540
 Gly Thr Trp Lys Arg Glu His Ser Lys His Ser Ser Ser Leu Pro Arg
 2545 2550 2555 2560
 Val Ser Thr Trp Arg Arg Thr Gly Ser Ser Ser Ser Ile Leu Ser Ala
 2565 2570 2575
 Ser Ser Glu Ser Ser Glu Lys Ala Lys Ser Glu Asp Glu Lys His Val
 2580 2585 2590
 Asn Ser Ile Ser Gly Thr Lys Gln Ser Lys Glu Asn Gln Val Ser Ala
 2595 2600 2605
 Lys Gly Thr Trp Arg Lys Ile Lys Glu Asn Glu Phe Ser Pro Thr Asn
 2610 2615 2620
 Ser Thr Ser Gln Thr Val Ser Ser Gly Ala Thr Asn Gly Ala Glu Ser
 2625 2630 2635 2640
 Lys Thr Leu Ile Tyr Gln Met Ala Pro Ala Val Ser Lys Thr Glu Asp
 2645 2650 2655
 Val Trp Val Arg Ile Glu Asp Cys Pro Ile Asn Asn Pro Arg Ser Gly
 2660 2665 2670
 Arg Ser Pro Thr Gly Asn Thr Pro Pro Val Ile Asp Ser Val Ser Glu
 2675 2680 2685
 Lys Ala Asn Pro Asn Ile Lys Asp Ser Lys Asp Asn Gln Ala Lys Gln

2690 2695 2700
 Asn Val Gly Asn Gly Ser Val Pro Met Arg Thr Val Gly Leu Glu Asn
 2705 2710 2715 2720
 Arg Leu Thr Ser Phe Ile Gln Val Asp Ala Pro Asp Gln Lys Gly Thr
 2725 2730 2735
 Glu Ile Lys Pro Gly Gln Asn Asn Pro Val Pro Val Ser Glu Thr Asn
 2740 2745 2750
 Glu Ser Pro Ile Val Glu Arg Thr Pro Phe Ser Ser Ser Ser Ser Ser
 2755 2760 2765
 Lys His Ser Ser Pro Ser Gly Thr Val Ala Ala Arg Val Thr Pro Phe
 2770 2775 2780
 Asn Tyr Asn Pro Ser Pro Arg Lys Ser Ser Ala Asp Ser Thr Ser Ala
 2785 2790 2795 2800
 Arg Pro Ser Gln Ile Pro Thr Pro Val Asn Asn Asn Thr Lys Lys Arg
 2805 2810 2815
 Asp Ser Lys Thr Asp Ser Thr Glu Ser Ser Gly Thr Gln Ser Pro Lys
 2820 2825 2830
 Arg His Ser Gly Ser Tyr Leu Val Thr Ser Val
 2835 2840

<210> 3
 <211> 3172
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(3172)
 <223> n = A,T,C or G

<400> 3
 gcagtcgccg ctccagtccta tccggcacta ggaacagccc cggngggcga gacgggtcccc 60
 gccatgtctg cggccatgag ggagaggttc gaccggttcc tgcacgagaa gaactgcatg 120
 actgaccttc tggccaagct cgaggccaaa accggcggtga acaggagctt catcgctctt 180
 ggtgtcatcg gactgggtggc cttgtacctg gtgttcgggt atggagcctc tctcctctgc 240
 aacctgatag gatttggcta ccagcctac atctcaatta aagctataga gagtcccaac 300
 aaagaagatg ataccagtg gctgacctac tgggtagtgt atggtgtgtt cagcattgct 360
 gaattcttct ctgatatctt cctgtcatgg tcccccttct actacatgct gaagtgtggc 420
 ttctgttgtt ggtgcatggc cccgagccct tctaattgggg ctgaactgct ctacaagcgc 480
 atcatccgtc ctttcttctt gaagcacgag tcccagatgg acagtgtggt caaggacctt 540
 aaagacaagt ccaaagagac tgcagatgcc atcactaaag aagcgaagaa agctaccgtg 600
 aattttactg gtgaagaaaa gaagagcacc taaaccagac taaaccagac tggatggaaa 660
 ctctctgccc tctctgtacc ttctactgg agcttgatgt tatattaggg actgtggtat 720
 aattatttta ataagtgtgc cttggaaaca tttttgagat attaaagatt ggaatgtgtt 780
 gtaagtctt ctgttactt ttactgtcta tatatatagg gaggacttta aacttaatgc 840
 agtgggcagt gtccacgttt ttggaaaatg tattttgcct ctgggtagga aaagatgtat 900
 gttgctatcc tgcaggaaat ataaacttaa aataaaaatta tatacccccac aggtgtgtga 960
 ctttactggg ctctccctgc acgsattttc tctgtagtta catttaggrt aatctttatg 1020
 gttctacttc ctrtaatgta caattttata taattcngra atgtttttaa tgtatttgtg 1080
 cacatgtaca tatggaaatg ttactgtctg actacancat gcatcatgct catggggagg 1140
 gagcagggga aggttgtatg tgtcatttat aacttctgta cagtaagacc acctgccaaa 1200
 agctggagga accattgtgc tgggtgtggc tactaaataa tacttttagga aatacgtgat 1260
 taatatgcaa gtgaacaaag tgagaaatga aatcgaatgg agattggcct gggtgtttcc 1320
 gtagtatatg gcatatgaat accaggatag ctttataaag cagttagtta gttagttact 1380
 cactctagtg ataaatcggg aaattttacac acacacacac acacacacac acacacacac 1440

acacacacac acacacacag agtaccctgt aactctcaat tccctgaaaa actagtaata 1500
 ctgtcttatt tgctataaac ttacatatt tgtctattgt caagatgcta cantggamnc 1560
 cttttctggg tttatcttca nagsggagan acatgttgat ttagtcttct ttcccaatct 1620
 tcttttttaa mccagtttna ggmncctctg ragatttgyt cacctctgat tacatgtatg 1680
 ttctygtttg tatcatkagc aacaacatgc taatgrcgac acctagctct ragmgcaatt 1740
 ctgggagant garaggnwgt ccataatctg cttggcaata gttaagtaa 1800
 tctatcttca gtttttctct ggcttttaag gtcaaacaca agaggcttcc ctagtttaca 1860
 agtcagagtc actttagatc catttaaagt cctcatccg tttctttgt gttgataagc 1920
 tgcacakgac tacatagtaa gtacaganca gtaaagttaa nncggatgtc tccattgatc 1980
 tgccaantcg ntatagagag caatttgtct ggactagaaa atctgagttt tacaccatac 2040
 ttttaagagt ccttttgaat taaactagac taaaacaagt gtataactaa actaacaaga 2100
 ttaaatatcc agccagtaca gtatttttta aggcaaataa agatgattag ctcaccttga 2160
 gntaacaatc aggttaagtc atnacaatgt ctcattgatgt naanaatatt aaagatatca 2220
 atactaagtg acagtatcac nnctaataata atatggatca gagcatttat tttggggagg 2280
 aaaacagtg tgattaccgg cattttatta aacttaaaac tttgtagaaa gcaaacaaaa 2340
 ttgttcttgg gagaaaatca acttttagat taaaaaaatt ttaagtawct aggagtattt 2400
 aaatcctttt ccataaata aaagtacagt tttcttgggt gcagaatgaa aatcagcaac 2460
 ntctagcata tagactatat aatcagattg acagcatata gaatatatta tcagacaaga 2520
 tgaggaggta caaaagttac tattgtctcat aatgacttac aggctaaaaan tagntntaaa 2580
 atactatatt aaattctgaa tgcaattttt ttttgttccc ttgagaccaa aatttaagtt 2640
 aactgttgcg ggcagtctaa gtgtaaatgt taacagcagg agaagttaa aattgagcag 2700
 ttctgtttgca tgatttccca aatgaaatac tgccttggct agagtttgaa aaactaattg 2760
 agcctgtgcc tggctagaaa acaagcgttt atttgaatgt gaatagtgtt tcaaaggtat 2820
 gtagttacag aattcctacc aaacagctta aattcttcaa gaaagaattc ctgcagcagt 2880
 tattccctta cctgaaggct tcaatcattt ggatcaacaa ctgctactct cggaagact 2940
 cctctactca cagctgaaga aaatgagcac acccttcaca ctgttatcac ctatcctgaa 3000
 gatgtgatac actgaatgga aataaataga tgtaaataaa attgagwtct catttaaaaa 3060
 aaaccatgtg cccaatggga aaatgacctc atgttgtggt ttaaacagca actgcacca 3120
 ctgacacagc ccattgagct ancctatata tacatctctg tcagtgcacc tc 3172

<210> 4
 <211> 210
 <212> PRT
 <213> Homo sapiens

<400> 4
 Ala Val Ala Ala Pro Val Tyr Pro Ala Leu Gly Thr Ala Pro Gly Gly
 1 5 10 15
 Glu Thr Val Pro Ala Met Ser Ala Ala Met Arg Glu Arg Phe Asp Arg
 20 25 30
 Phe Leu His Glu Lys Asn Cys Met Thr Asp Leu Leu Ala Lys Leu Glu
 35 40 45
 Ala Lys Thr Gly Val Asn Arg Ser Phe Ile Ala Leu Gly Val Ile Gly
 50 55 60
 Leu Val Ala Leu Tyr Leu Val Phe Gly Tyr Gly Ala Ser Leu Leu Cys
 65 70 75 80
 Asn Leu Ile Gly Phe Gly Tyr Pro Ala Tyr Ile Ser Ile Lys Ala Ile
 85 90 95
 Glu Ser Pro Asn Lys Glu Asp Asp Thr Gln Trp Leu Thr Tyr Trp Val
 100 105 110
 Val Tyr Gly Val Phe Ser Ile Ala Glu Phe Phe Ser Asp Ile Phe Leu
 115 120 125
 Ser Trp Phe Pro Phe Tyr Tyr Met Leu Lys Cys Gly Phe Leu Leu Trp
 130 135 140
 Cys Met Ala Pro Ser Pro Ser Asn Gly Ala Glu Leu Leu Tyr Lys Arg

145
 Ile Ile Arg Pro Phe Phe Leu Lys His Glu Ser Gln Met Asp Ser Val
 150
 165
 170
 Val Lys Asp Leu Lys Asp Lys Ser Lys Glu Thr Ala Asp Ala Ile Thr
 180
 185
 Lys Glu Ala Lys Lys Ala Thr Val Asn Leu Leu Gly Glu Glu Lys Lys
 195
 200
 205
 Ser Thr
 210

<210> 5
 <211> 434
 <212> PRT
 <213> Homo sapiens

<400> 5
 Val Ala Pro Val Val Val Gly Ser Gly Arg Ala Pro Arg His Pro Ala
 1
 5
 10
 Pro Ala Ala Met His Pro Arg Arg Pro Asp Gly Phe Asp Gly Leu Gly
 20
 25
 30
 Tyr Arg Gly Gly Ala Arg Asp Glu Gln Gly Phe Gly Gly Ala Phe Pro
 35
 40
 45
 Ala Arg Ser Phe Ser Thr Gly Ser Asp Leu Gly His Trp Val Thr Thr
 50
 55
 60
 Pro Pro Asp Ile Pro Gly Ser Arg Asn Leu His Trp Gly Glu Lys Ser
 65
 70
 75
 Pro Pro Tyr Gly Val Pro Thr Thr Ser Thr Pro Tyr Glu Gly Pro Thr
 85
 90
 95
 Glu Glu Pro Phe Ser Ser Gly Gly Gly Ser Val Gln Gly Gln Ser
 100
 105
 110
 Ser Glu Gln Leu Asn Arg Phe Ala Gly Phe Gly Ile Gly Leu Ala Ser
 115
 120
 125
 Leu Phe Thr Glu Asn Val Leu Ala His Pro Cys Ile Val Leu Arg Arg
 130
 135
 140
 Gln Cys Gln Val Asn Tyr His Ala Gln His Tyr His Leu Thr Pro Phe
 145
 150
 155
 Thr Val Ile Asn Ile Met Tyr Ser Phe Asn Lys Thr Gln Gly Pro Arg
 165
 170
 175
 Ala Leu Trp Lys Gly Met Gly Ser Thr Phe Ile Val Gln Gly Val Thr
 180
 185
 190
 Leu Gly Ala Glu Gly Ile Ile Ser Glu Phe Thr Pro Leu Pro Arg Glu
 195
 200
 205
 Val Leu His Lys Trp Ser Pro Lys Gln Ile Gly Glu His Leu Leu Leu
 210
 215
 220
 Lys Ser Leu Thr Tyr Val Val Ala Met Pro Phe Tyr Ser Ala Ser Leu
 225
 230
 235
 Ile Glu Thr Val Gln Ser Glu Ile Ile Arg Asp Asn Thr Gly Ile Leu
 245
 250
 255
 Glu Cys Val Lys Glu Gly Ile Gly Arg Val Ile Gly Met Gly Val Pro
 260
 265
 270
 His Ser Lys Arg Leu Leu Pro Leu Ser Leu Ile Phe Pro Thr Val
 275
 280
 285
 Leu His Gly Val Leu His Tyr Ile Ile Ser Ser Val Ile Gln Lys Phe
 290
 295
 300
 Val Leu Leu Ile Leu Lys Arg Lys Thr Tyr Asn Ser His Leu Ala Glu

305 310 315 320
 Ser Thr Ser Pro Val Gln Ser Met Leu Asp Ala Tyr Phe Pro Glu Leu
 325 330 335
 Ile Ala Asn Phe Ala Ala Ser Leu Cys Ser Asp Val Ile Leu Tyr Pro
 340 345 350
 Leu Glu Thr Val Leu His Arg Leu His Ile Gln Gly Thr Arg Thr Ile
 355 360 365
 Ile Asp Asn Thr Asp Leu Gly Tyr Glu Val Leu Pro Ile Asn Thr Gln
 370 375 380
 Tyr Glu Gly Met Arg Asp Cys Ile Asn Thr Ile Arg Gln Glu Glu Gly
 385 390 395 400
 Val Phe Gly Phe Tyr Lys Gly Phe Gly Ala Val Ile Ile Gln Tyr Thr
 405 410 415
 Leu His Ala Ala Val Leu Gln Ile Thr Lys Ile Ile Tyr Ser Thr Leu
 420 425 430
 Leu Gln

<210> 6
 <211> 185
 <212> PRT
 <213> Homo sapiens

<400> 6
 Glu Leu Arg Arg Phe Asp Arg Phe Leu His Glu Lys Asn Cys Met Thr
 1 5 10 15
 Asp Leu Leu Ala Lys Leu Glu Ala Lys Thr Gly Val Asn Arg Ser Phe
 20 25 30
 Ile Ala Leu Gly Val Ile Gly Leu Val Ala Leu Tyr Leu Val Phe Gly
 35 40 45
 Tyr Gly Ala Ser Leu Leu Cys Asn Leu Ile Gly Phe Gly Tyr Pro Ala
 50 55 60
 Tyr Ile Ser Ile Lys Ala Ile Glu Ser Pro Asn Lys Glu Asp Asp Thr
 65 70 75 80
 Gln Trp Leu Thr Tyr Trp Val Val Tyr Gly Val Phe Ser Ile Ala Glu
 85 90 95
 Phe Phe Ser Asp Ile Phe Leu Ser Trp Phe Pro Phe Tyr Tyr Ile Leu
 100 105 110
 Lys Cys Gly Phe Leu Leu Trp Cys Met Ala Pro Ser Pro Ser Asn Gly
 115 120 125
 Ala Glu Leu Leu Tyr Lys Arg Ile Ile Arg Pro Phe Phe Leu Lys His
 130 135 140
 Glu Ser Gln Met Asp Ser Val Val Lys Asp Leu Lys Asp Lys Ala Lys
 145 150 155 160
 Glu Thr Ala Asp Ala Ile Thr Lys Glu Ala Lys Lys Ala Thr Val Asn
 165 170 175
 Leu Leu Gly Glu Glu Lys Lys Ser Thr
 180 185

<210> 7
 <211> 2842
 <212> PRT
 <213> Homo sapiens

<400> 7

Met Ala Ala Ala Ser Tyr Asp Gln Leu Leu Lys Gln Val Glu Ala Leu
 1 5 10 15
 Lys Met Glu Asn Ser Asn Leu Arg Gln Glu Leu Glu Asp Asn Ser Asn
 20 25 30
 His Leu Thr Lys Leu Glu Thr Glu Ala Ser Asn Met Lys Glu Val Leu
 35 40 45
 Lys Gln Leu Gln Gly Ser Ile Glu Asp Glu Ala Met Lys Glu Val Leu
 50 55 60
 Gln Ile Asp Leu Leu Glu Arg Leu Lys Glu Leu Asn Leu Asp Ser Ser
 65 70 75 80
 Asn Phe Pro Gly Val Lys Leu Arg Ser Lys Met Ser Leu Arg Ser Tyr
 85 90 95
 Gly Ser Arg Glu Gly Ser Val Ser Ser Arg Ser Gly Glu Cys Ser Pro
 100 105 110
 Val Pro Met Gly Ser Phe Pro Arg Arg Gly Phe Val Asn Gly Ser Arg
 115 120 125
 Glu Ser Thr Gly Tyr Leu Glu Glu Leu Glu Lys Glu Arg Ser Leu Leu
 130 135 140
 Leu Ala Asp Leu Asp Lys Glu Glu Lys Glu Lys Asp Trp Tyr Tyr Ala
 145 150 155 160
 Gln Leu Gln Asn Leu Thr Lys Arg Ile Asp Ser Leu Pro Thr Glu Asn
 165 170 175
 Phe Ser Leu Gln Thr Asp Met Thr Arg Gln Leu Glu Tyr Glu Ala
 180 185 190
 Arg Gln Ile Arg Val Ala Met Glu Glu Gln Leu Gly Thr Cys Gln Asp
 195 200 205
 Met Glu Lys Arg Ala Gln Arg Arg Ile Ala Arg Ile Gln Gln Ile Glu
 210 215 220
 Lys Asp Ile Leu Arg Ile Arg Gln Leu Leu Gln Ser Gln Ala Thr Glu
 225 230 235 240
 Ala Glu Arg Ser Ser Gln Asn Lys His Glu Thr Gly Ser His Asp Ala
 245 250 255
 Glu Arg Gln Asn Glu Gly Gln Gly Val Gly Glu Ile Asn Met Ala Thr
 260 265 270
 Ser Gly Asn Gly Gln Gly Ser Thr Arg Met Asp His Glu Thr Ala
 275 280 285
 Ser Val Leu Ser Ser Ser Ser Thr His Ser Ala Pro Arg Arg Leu Thr
 290 295 300
 Ser His Leu Gly Thr Lys Val Glu Met Val Tyr Ser Leu Leu Ser Met
 305 310 315 320
 Leu Gly Thr His Asp Lys Asp Asp Met Ser Arg Thr Leu Leu Ala Met
 325 330 335
 Ser Ser Ser Gln Asp Ser Cys Ile Ser Met Arg Gln Ser Gly Cys Leu
 340 345 350
 Pro Leu Leu Ile Gln Leu Leu His Gly Asn Asp Lys Asp Ser Val Leu
 355 360 365
 Leu Gly Asn Ser Arg Gly Ser Lys Glu Ala Arg Ala Arg Ala Ser Ala
 370 375 380
 Ala Leu His Asn Ile Ile His Ser Gln Pro Asp Asp Lys Arg Gly Arg
 385 390 395 400
 Arg Glu Ile Arg Val Leu His Leu Leu Glu Gln Ile Arg Ala Tyr Cys
 405 410 415
 Glu Thr Cys Trp Glu Trp Gln Glu Ala His Glu Pro Gly Met Asp Gln
 420 425 430
 Asp Lys Asn Pro Met Pro Ala Pro Val Glu His Gln Ile Cys Pro Ala

435 440 445
 Val Cys Val Leu Met Lys Leu Ser Phe Asp Glu Glu His Arg His Ala
 450 455 460
 Met Asn Glu Leu Gly Gly Leu Gln Ala Ile Ala Glu Leu Leu Gln Val
 465 470 475 480
 Asp Cys Glu Met Tyr Gly Leu Thr Asn Asp His Tyr Ser Ile Thr Leu
 485 490 495
 Arg Arg Tyr Ala Gly Met Ala Leu Thr Asn Leu Thr Phe Gly Asp Val
 500 505 510
 Ala Asn Lys Ala Thr Leu Cys Ser Met Lys Gly Cys Met Arg Ala Leu
 515 520 525
 Val Ala Gln Leu Lys Ser Glu Ser Glu Asp Leu Gln Gln Val Ile Ala
 530 535 540
 Ser Val Leu Arg Asn Leu Ser Trp Arg Ala Asp Val Asn Ser Lys Lys
 545 550 555 560
 Thr Leu Arg Glu Val Gly Ser Val Lys Ala Leu Met Glu Cys Ala Leu
 565 570 575
 Glu Val Lys Lys Glu Ser Thr Leu Lys Ser Val Leu Ser Ala Leu Trp
 580 585 590
 Asn Leu Ser Ala His Cys Thr Glu Asn Lys Ala Asp Ile Cys Ala Val
 595 600 605
 Asp Gly Ala Leu Ala Phe Leu Val Gly Thr Leu Thr Tyr Arg Ser Gln
 610 615 620
 Thr Asn Thr Leu Ala Ile Ile Glu Ser Gly Gly Gly Ile Leu Arg Asn
 625 630 635 640
 Val Ser Ser Leu Ile Ala Thr Asn Glu Asp His Arg Gln Ile Leu Arg
 645 650 655
 Glu Asn Asn Cys Leu Gln Thr Leu Leu Gln His Leu Lys Ser His Ser
 660 665 670
 Leu Thr Ile Val Ser Asn Ala Cys Gly Thr Leu Trp Asn Leu Ser Ala
 675 680 685
 Arg Asn Pro Lys Asp Gln Glu Ala Leu Trp Asp Met Gly Ala Val Ser
 690 695 700
 Met Leu Lys Asn Leu Ile His Ser Lys His Lys Met Ile Ala Met Gly
 705 710 715 720
 Ser Ala Ala Ala Leu Arg Asn Leu Met Ala Asn Arg Pro Ala Lys Tyr
 725 730 735
 Lys Asp Ala Asn Ile Met Ser Pro Gly Ser Ser Leu Pro Ser Leu His
 740 745 750
 Val Arg Lys Gln Lys Ala Leu Glu Ala Glu Leu Asp Ala Gln His Leu
 755 760 765
 Ser Glu Thr Phe Asp Asn Ile Asp Asn Leu Ser Pro Lys Ala Ser His
 770 775 780
 Arg Ser Lys Gln Arg His Lys Gln Ser Leu Tyr Gly Asp Tyr Val Phe
 785 790 795 800
 Asp Thr Asn Arg His Asp Asp Asn Arg Ser Asp Asn Phe Asn Thr Gly
 805 810 815
 Asn Met Thr Val Leu Ser Pro Tyr Leu Asn Thr Thr Val Leu Pro Ser
 820 825 830
 Ser Ser Ser Ser Arg Gly Ser Leu Asp Ser Ser Arg Ser Glu Lys Asp
 835 840 845
 Arg Ser Leu Glu Arg Glu Arg Gly Ile Gly Leu Gly Asn Tyr His Pro
 850 855 860
 Ala Thr Glu Asn Pro Gly Thr Ser Ser Lys Arg Gly Leu Gln Ile Ser
 865 870 875 880

Thr Thr Ala Ala Gln Ile Ala Lys Val Met Glu Glu Val Ser Ala Ile
 885 890 895
 His Thr Ser Gln Glu Asp Arg Ser Ser Gly Ser Thr Thr Glu Leu His
 900 905 910
 Cys Val Thr Asp Glu Arg Asn Ala Leu Arg Arg Ser Ser Ala Ala His
 915 920 925
 Thr His Ser Asn Thr Tyr Asn Phe Thr Lys Ser Glu Asn Ser Asn Arg
 930 935 940
 Thr Cys Ser Met Pro Tyr Ala Lys Leu Glu Tyr Lys Arg Ser Ser Asn
 945 950 955 960
 Asp Ser Leu Asn Ser Val Ser Ser Ser Asp Gly Tyr Gly Lys Arg Gly
 965 970 975
 Gln Met Lys Pro Ser Ile Glu Ser Tyr Ser Glu Asp Asp Glu Ser Lys
 980 985 990
 Phe Cys Ser Tyr Gly Gln Tyr Pro Ala Asp Leu Ala His Lys Ile His
 995 1000 1005
 Ser Ala Asn His Met Asp Asp Asn Asp Gly Glu Leu Asp Thr Pro Ile
 1010 1015 1020
 Asn Tyr Ser Leu Lys Tyr Ser Asp Glu Gln Leu Asn Ser Gly Arg Gln
 1025 1030 1035 1040
 Ser Pro Ser Gln Asn Glu Arg Trp Ala Arg Pro Lys His Ile Ile Glu
 1045 1050 1055
 Asp Glu Ile Lys Gln Ser Glu Gln Arg Gln Ser Arg Asn Gln Ser Thr
 1060 1065 1070
 Thr Tyr Pro Val Tyr Thr Glu Ser Thr Asp Asp Lys His Leu Lys Phe
 1075 1080 1085
 Gln Pro His Phe Gly Gln Gln Glu Cys Val Ser Pro Tyr Arg Ser Arg
 1090 1095 1100
 Gly Ala Asn Gly Ser Glu Thr Asn Arg Val Gly Ser Asn His Gly Ile
 1105 1110 1115 1120
 Asn Gln Asn Val Ser Gln Ser Leu Cys Gln Glu Asp Asp Tyr Glu Asp
 1125 1130 1135
 Asp Lys Pro Thr Asn Tyr Ser Glu Arg Tyr Ser Glu Glu Glu Gln His
 1140 1145 1150
 Glu Glu Glu Glu Arg Pro Thr Asn Tyr Ser Ile Lys Tyr Asn Glu Glu
 1155 1160 1165
 Lys Arg His Val Asp Gln Pro Ile Asp Tyr Ser Leu Lys Tyr Ala Thr
 1170 1175 1180
 Asp Ile Pro Ser Ser Gln Lys Gln Ser Phe Ser Phe Ser Lys Ser Ser
 1185 1190 1195 1200
 Ser Gly Gln Ser Ser Lys Thr Glu His Met Ser Ser Ser Ser Glu Asn
 1205 1210 1215
 Thr Ser Thr Pro Ser Ser Asn Ala Lys Arg Gln Asn Gln Leu His Pro
 1220 1225 1230
 Ser Ser Ala Gln Ser Arg Ser Gly Gln Pro Gln Lys Ala Ala Thr Cys
 1235 1240 1245
 Lys Val Ser Ser Ile Asn Gln Glu Thr Ile Gln Thr Tyr Cys Val Glu
 1250 1255 1260
 Asp Thr Pro Ile Cys Phe Ser Arg Cys Ser Ser Leu Ser Ser Leu Ser
 1265 1270 1275 1280
 Ser Ala Glu Asp Glu Ile Gly Cys Asn Gln Thr Thr Gln Glu Ala Asp
 1285 1290 1295
 Ser Ala Asn Thr Leu Gln Ile Ala Glu Ile Lys Glu Lys Ile Gly Thr
 1300 1305 1310
 Arg Ser Ala Glu Asp Pro Val Ser Glu Val Pro Ala Val Ser Gln His

1315 1320 1325
 Pro Arg Thr Lys Ser Ser Arg Leu Gln Gly Ser Ser Leu Ser Ser Glu
 1330 1335 1340
 Ser Ala Arg His Lys Ala Val Glu Phe Ser Ser Gly Ala Lys Ser Pro
 1345 1350 1355 1360
 Ser Lys Ser Gly Ala Gln Thr Pro Lys Ser Pro Pro Glu His Tyr Val
 1365 1370 1375
 Gln Glu Thr Pro Leu Met Phe Ser Arg Cys Thr Ser Val Ser Ser Leu
 1380 1385 1390
 Asp Ser Phe Glu Ser Arg Ser Ile Ala Ser Ser Val Gln Ser Glu Pro
 1395 1400 1405
 Cys Ser Gly Met Val Ser Gly Ile Ile Ser Pro Ser Asp Leu Pro Asp
 1410 1415 1420
 Ser Pro Gly Gln Thr Met Pro Pro Ser Arg Ser Lys Thr Pro Pro Pro
 1425 1430 1435 1440
 Pro Pro Gln Thr Ala Gln Thr Lys Arg Glu Val Pro Lys Asn Lys Ala
 1445 1450 1455
 Pro Thr Ala Glu Lys Arg Glu Ser Gly Pro Lys Gln Ala Ala Val Asn
 1460 1465 1470
 Ala Ala Val Gln Arg Val Gln Val Leu Pro Asp Ala Asp Thr Leu Leu
 1475 1480 1485
 His Phe Ala Thr Glu Ser Thr Pro Asp Gly Phe Ser Cys Ser Ser Ser
 1490 1495 1500
 Leu Ser Ala Leu Ser Leu Asp Glu Pro Phe Ile Gln Lys Asp Val Glu
 1505 1510 1515 1520
 Leu Arg Ile Met Pro Pro Val Gln Glu Asn Asp Asn Gly Asn Glu Thr
 1525 1530 1535
 Glu Ser Glu Gln Pro Lys Glu Ser Asn Glu Asn Gln Glu Lys Glu Ala
 1540 1545 1550
 Glu Lys Thr Ile Asp Ser Glu Lys Asp Leu Leu Asp Asp Ser Asp Asp
 1555 1560 1565
 Asp Asp Ile Glu Ile Leu Glu Glu Cys Ile Ile Ser Ala Met Pro Thr
 1570 1575 1580
 Lys Ser Ser Arg Lys Ala Lys Lys Pro Ala Gln Thr Ala Ser Lys Leu
 1585 1590 1595 1600
 Pro Pro Pro Val Ala Arg Lys Pro Ser Gln Leu Pro Val Tyr Lys Leu
 1605 1610 1615
 Leu Pro Ser Gln Asn Arg Leu Gln Pro Gln Lys His Val Ser Phe Thr
 1620 1625 1630
 Pro Gly Asp Asp Met Pro Arg Val Tyr Cys Val Glu Gly Thr Pro Ile
 1635 1640 1645
 Asn Phe Ser Thr Ala Thr Ser Leu Ser Asp Leu Thr Ile Glu Ser Pro
 1650 1655 1660
 Pro Asn Glu Leu Ala Ala Gly Glu Gly Val Arg Gly Gly Ala Gln Ser
 1665 1670 1675 1680
 Gly Glu Phe Glu Lys Arg Asp Thr Ile Pro Thr Glu Gly Arg Ser Thr
 1685 1690 1695
 Asp Glu Ala Gln Gly Gly Lys Thr Ser Ser Val Thr Ile Pro Glu Leu
 1700 1705 1710
 Asp Asp Asn Lys Ala Glu Glu Gly Asp Ile Leu Ala Glu Cys Ile Asn
 1715 1720 1725
 Ser Ala Met Pro Lys Gly Lys Ser His Lys Pro Phe Arg Val Lys Lys
 1730 1735 1740
 Ile Met Asp Gln Val Gln Gln Ala Ser Ala Ser Ser Ser Ala Pro Asn
 1745 1750 1755 1760

Lys Asn Gln Leu Asp Gly Lys Lys Lys Lys Pro Thr Ser Pro Val Lys
 1765 1770 1775
 Pro Ile Pro Gln Asn Thr Glu Tyr Arg Thr Arg Val Arg Lys Asn Ala
 1780 1785 1790
 Asp Ser Lys Asn Asn Leu Asn Ala Glu Arg Val Phe Ser Asp Asn Lys
 1795 1800 1805
 Asp Ser Lys Lys Gln Asn Leu Lys Asn Asn Ser Lys Asp Phe Asn Asp
 1810 1815 1820
 Lys Leu Pro Asn Asn Glu Asp Arg Val Arg Gly Ser Phe Ala Phe Asp
 1825 1830 1835 1840
 Ser Pro His His Tyr Thr Pro Ile Glu Gly Thr Pro Tyr Cys Phe Ser
 1845 1850 1855
 Arg Asn Asp Ser Leu Ser Ser Leu Asp Phe Asp Asp Asp Val Asp
 1860 1865 1870
 Leu Ser Arg Glu Lys Ala Glu Leu Arg Lys Ala Lys Glu Asn Lys Glu
 1875 1880 1885
 Ser Glu Ala Lys Val Thr Ser His Thr Glu Leu Thr Ser Asn Gln Gln
 1890 1895 1900
 Ser Ala Asn Lys Thr Gln Ala Ile Ala Lys Gln Pro Ile Asn Arg Gly
 1905 1910 1915 1920
 Gln Pro Lys Pro Ile Leu Gln Lys Gln Ser Thr Phe Pro Gln Ser Ser
 1925 1930 1935
 Lys Asp Ile Pro Asp Arg Gly Ala Ala Thr Asp Glu Lys Leu Gln Asn
 1940 1945 1950
 Phe Ala Ile Glu Asn Thr Pro Val Cys Phe Ser His Asn Ser Ser Leu
 1955 1960 1965
 Ser Ser Leu Ser Asp Ile Asp Gln Glu Asn Asn Asn Lys Glu Asn Glu
 1970 1975 1980
 Pro Ile Lys Glu Thr Glu Pro Pro Asp Ser Gln Gly Glu Pro Ser Lys
 1985 1990 1995 2000
 Pro Gln Ala Ser Gly Tyr Ala Pro Lys Ser Phe His Val Glu Asp Thr
 2005 2010 2015
 Pro Val Cys Phe Ser Arg Asn Ser Ser Leu Ser Ser Leu Ser Ile Asp
 2020 2025 2030
 Ser Glu Asp Asp Leu Leu Gln Glu Cys Ile Ser Ser Ala Met Pro Lys
 2035 2040 2045
 Lys Lys Lys Pro Ser Arg Leu Lys Gly Asp Asn Glu Lys His Ser Pro
 2050 2055 2060
 Arg Asn Met Gly Gly Ile Leu Gly Glu Asp Leu Thr Leu Asp Leu Lys
 2065 2070 2075 2080
 Asp Ile Gln Arg Pro Asp Ser Glu His Gly Leu Ser Pro Asp Ser Glu
 2085 2090 2095
 Asn Phe Asp Trp Lys Ala Ile Gln Glu Gly Ala Asn Ser Ile Val Ser
 2100 2105 2110
 Ser Leu His Gln Ala Ala Ala Ala Cys Leu Ser Arg Gln Ala Ser
 2115 2120 2125
 Ser Asp Ser Asp Ser Ile Leu Ser Leu Lys Ser Gly Ile Ser Leu Gly
 2130 2135 2140
 Ser Pro Phe His Leu Thr Pro Asp Gln Glu Glu Lys Pro Phe Thr Ser
 2145 2150 2155 2160
 Asn Lys Gly Pro Arg Ile Leu Lys Pro Gly Glu Lys Ser Thr Leu Glu
 2165 2170 2175
 Thr Lys Lys Ile Glu Ser Glu Ser Lys Gly Ile Lys Gly Lys Lys
 2180 2185 2190
 Val Tyr Lys Ser Leu Ile Thr Gly Lys Val Arg Ser Asn Ser Glu Ile

2195 2200 2205
 Ser Gly Gln Met Lys Gln Pro Leu Gln Ala Asn Met Pro Ser Ile Ser
 2210 2215 2220
 Arg Gly Arg Thr Met Ile His Ile Pro Gly Val Arg Asn Ser Ser Ser
 2225 2230 2235 2240
 Ser Thr Ser Pro Val Ser Lys Lys Gly Pro Pro Leu Lys Thr Pro Ala
 2245 2250 2255
 Ser Lys Ser Pro Ser Glu Gly Gln Thr Ala Thr Thr Ser Pro Arg Gly
 2260 2265 2270
 Ala Lys Pro Ser Val Lys Ser Glu Leu Ser Pro Val Ala Arg Gln Thr
 2275 2280 2285
 Ser Gln Ile Gly Gly Ser Ser Lys Ala Pro Ser Arg Ser Gly Ser Arg
 2290 2295 2300
 Asp Ser Thr Pro Ser Arg Pro Ala Gln Gln Pro Leu Ser Arg Pro Ile
 2305 2310 2315 2320
 Gln Ser Pro Gly Arg Asn Ser Ile Ser Pro Gly Arg Asn Gly Ile Ser
 2325 2330 2335
 Pro Pro Asn Lys Leu Ser Gln Leu Pro Arg Thr Ser Ser Pro Ser Thr
 2340 2345 2350
 Ala Ser Thr Lys Ser Ser Gly Ser Gly Lys Met Ser Tyr Thr Ser Pro
 2355 2360 2365
 Gly Arg Gln Met Ser Gln Gln Asn Leu Thr Lys Gln Thr Gly Leu Ser
 2370 2375 2380
 Lys Asn Ala Ser Ser Ile Pro Arg Ser Glu Ser Ala Ser Lys Gly Leu
 2385 2390 2395 2400
 Asn Gln Met Asn Asn Gly Asn Gly Ala Asn Lys Lys Val Glu Leu Ser
 2405 2410 2415
 Arg Met Ser Ser Thr Lys Ser Ser Gly Ser Glu Ser Asp Arg Ser Glu
 2420 2425 2430
 Arg Pro Val Leu Val Arg Gln Ser Thr Phe Ile Lys Glu Ala Pro Ser
 2435 2440 2445
 Pro Thr Leu Arg Arg Lys Leu Glu Glu Ser Ala Ser Phe Glu Ser Leu
 2450 2455 2460
 Ser Pro Ser Ser Arg Pro Ala Ser Pro Thr Arg Ser Gln Ala Gln Thr
 2465 2470 2475 2480
 Pro Val Leu Ser Pro Ser Leu Pro Asp Met Ser Leu Ser Thr His Ser
 2485 2490 2495
 Ser Val Gln Ala Gly Gly Trp Arg Lys Leu Pro Pro Asn Leu Ser Pro
 2500 2505 2510
 Thr Ile Glu Tyr Asn Asp Gly Arg Pro Ala Lys Arg His Asp Ile Ala
 2515 2520 2525
 Arg Ser His Ser Glu Ser Pro Ser Arg Leu Pro Ile Asn Arg Ser Gly
 2530 2535 2540
 Thr Trp Lys Arg Glu His Ser Lys His Ser Ser Ser Leu Pro Arg Val
 2545 2550 2555 2560
 Ser Thr Trp Arg Arg Thr Gly Ser Ser Ser Ser Ile Leu Ser Ala Ser
 2565 2570 2575
 Ser Glu Ser Ser Glu Lys Ala Lys Ser Glu Asp Glu Lys His Val Asn
 2580 2585 2590
 Ser Ile Ser Gly Thr Lys Gln Ser Lys Glu Asn Gln Val Ser Ala Lys
 2595 2600 2605
 Gly Thr Trp Arg Lys Ile Lys Glu Asn Glu Phe Ser Pro Thr Asn Ser
 2610 2615 2620
 Thr Ser Gln Thr Val Ser Ser Gly Ala Thr Asn Gly Ala Glu Ser Lys
 2625 2630 2635 2640

Thr Leu Ile Tyr Gln Met Ala Pro Ala Val Ser Lys Thr Glu Asp Val
 2645 2650 2655
 Trp Val Arg Ile Glu Asp Cys Pro Ile Asn Asn Pro Arg Ser Gly Arg
 2660 2665 2670
 Ser Pro Thr Gly Asn Thr Pro Pro Val Ile Asp Ser Val Ser Glu Lys
 2675 2680 2685
 Ala Asn Pro Asn Ile Lys Asp Ser Lys Asp Asn Gln Ala Lys Gln Asn
 2690 2695 2700
 Val Gly Asn Gly Ser Val Pro Met Arg Thr Val Gly Leu Glu Asn Arg
 2705 2710 2715 2720
 Leu Asn Ser Phe Ile Gln Val Asp Ala Pro Asp Gln Lys Gly Thr Glu
 2725 2730 2735
 Ile Lys Pro Gly Gln Asn Asn Pro Val Pro Val Ser Glu Thr Asn Glu
 2740 2745 2750
 Ser Ser Ile Val Glu Arg Thr Pro Phe Ser Ser Ser Ser Ser Lys
 2755 2760 2765
 His Ser Ser Pro Ser Gly Thr Val Ala Ala Arg Val Thr Pro Phe Asn
 2770 2775 2780
 Tyr Asn Pro Ser Pro Arg Lys Ser Ser Ala Asp Ser Thr Ser Ala Arg
 2785 2790 2795 2800
 Pro Ser Gln Ile Pro Thr Pro Val Asn Asn Asn Thr Lys Lys Arg Asp
 2805 2810 2815
 Ser Lys Thr Asp Ser Thr Glu Ser Ser Gly Thr Gln Ser Pro Lys Arg
 2820 2825 2830
 His Ser Gly Ser Tyr Leu Val Thr Ser Val
 2835 2840

<210> 8
 <211> 31
 <212> PRT
 <213> Yeast

<400> 8
 Leu Thr Gly Ala Lys Gly Leu Gln Leu Arg Ala Leu Arg Arg Ile Ala
 1 5 10 15
 Arg Ile Glu Gln Gly Gly Thr Ala Ile Ser Pro Thr Ser Pro Leu
 20 25 30

<210> 9
 <211> 29
 <212> PRT
 <213> Homo sapiens

<400> 9
 Leu Tyr Trp Arg Ile Tyr Lys Glu Thr Glu Lys Arg Thr Lys Glu Leu
 1 5 10 15
 Ala Gly Leu Gln Ala Ser Gly Thr Glu Ala Glu Thr Glu
 20 25

<210> 10
 <211> 29
 <212> PRT
 <213> Homo sapiens

<400> 10

Leu Tyr Pro Asn Leu Ala Glu Glu Arg Ser Arg Trp Glu Lys Glu Leu
 1 5 10 15
 Ala Gly Leu Arg Glu Glu Asn Glu Ser Leu Thr Ala Met
 20 25

<210> 11
 <211> 40
 <212> DNA
 <213> Homo sapiens

<400> 11 40
 gtatcaagac tgtgactttt aattgtagtt tatccatttt

<210> 12
 <211> 40
 <212> DNA
 <213> Homo sapiens

<400> 12 40
 tttagaattt catgttaata tattgtgttc tttttaacag

<210> 13
 <211> 40
 <212> DNA
 <213> Homo sapiens

<400> 13 40
 gtagatttta aaaagggtgtt ttaaaataat tttttaagct

<210> 14
 <211> 40
 <212> DNA
 <213> Homo sapiens

<400> 14 40
 aagcaattgt tgtataaaaa cttgtttcta ttttatttag

<210> 15
 <211> 40
 <212> DNA
 <213> Homo sapiens

<400> 15 40
 gtaacttttc ttcatatagt aaacattgcc ttgtgtactc

<210> 16
 <211> 40
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(40)
 <223> n = A,T,C or G

<400> 16
nnnnnnnnnn nnngtccctt tttttaaaaa aaaaaaatag

40

<210> 17
<211> 40
<212> DNA
<213> Homo sapiens

<400> 17
gtaagtaact tggcagtaca acttatttga aactttaata

40

<210> 18
<211> 40
<212> DNA
<213> Homo sapiens

<400> 18
atacaagata ttgatacttt tttattattt gtgggttttag

40

<210> 19
<211> 40
<212> DNA
<213> Homo sapiens

<400> 19
gtaagttact tgtttctaag tgataaaaca gygaagagct

40

<210> 20
<211> 40
<212> DNA
<213> Homo sapiens

<400> 20
aataaaaaca taactaatta ggtttcttgt tttatttttag

40

<210> 21
<211> 40
<212> DNA
<213> Homo sapiens

<400> 21
gttagtaaat tscctttttt gtttgtgggt ataaaaatag

40

<210> 22
<211> 40
<212> DNA
<213> Homo sapiens

<400> 22
accatttttg catgtactga tggttaactcc atcttaacag

40

<210> 23
<211> 40
<212> DNA
<213> Homo sapiens

<400> 23 40
 gtaaataaat tattttatca tattttttaa aattatttaa

 <210> 24
 <211> 64
 <212> DNA
 <213> Homo sapiens

 <400> 24 60
 catgatgtta tctgtattta cctatagtct aaattatacc atctataatg tgcttaattt 64
 ttag

 <210> 25
 <211> 52
 <212> DNA
 <213> Homo sapiens

 <400> 25 52
 gtaacagaag attacaaacc ctggtcacta atgccatgac tactttgcta ag

 <210> 26
 <211> 46
 <212> DNA
 <213> Homo sapiens

 c64 <400> 26 46
 ggatattaaa gtcgtaattt tgtttctaaa ctcatttggc ccacag

 <210> 27
 <211> 40
 <212> DNA
 <213> Homo sapiens

 <400> 27 40
 gtatgttctc tatagtgtac atcgtagtgc atgtttcaaa

 <210> 28
 <211> 56
 <212> DNA
 <213> Homo sapiens

 <400> 28 56
 catcattgct cttcaaataa caaagcatta tggtttatgt tgattttatt tttcag

 <210> 29
 <211> 43
 <212> DNA
 <213> Homo sapiens

 <400> 29 43
 gtaagacaaa aatgtttttt aatgacatag acaattactg gtg

 <210> 30
 <211> 40

<212> DNA
<213> Homo sapiens

<400> 30
ttagatgatt gtctttttcc tcttgccctt tttaaattag

40

<210> 31
<211> 44
<212> DNA
<213> Homo sapiens

<400> 31
gtatgttttt ataacatgta tttcttaaga tagctcaggt atga

44

<210> 32
<211> 54
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(54)
<223> n = A,T,C or G

<400> 32
gcttggttc aagttgnctt tttaatgatc ctctattctg tatttaattt acag

54

<210> 33
<211> 65
<212> DNA
<213> Homo sapiens

<400> 33
gtactattta gaatttcacc tgtttttctt tttctctttt ttctttgagg cagggcttca
ctctg

60
65

<210> 34
<211> 52
<212> DNA
<213> Homo sapiens

<400> 34
gcaactagta tgattttatg tataaattaa tctaaaattg attaatttcc ag

52

<210> 35
<211> 42
<212> DNA
<213> Homo sapiens

<400> 35
gtaccttga aaacatttag tactataata tgaatttcat gt

42

<210> 36
<211> 40
<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(40)

<223> n = A,T,C or G

<400> 36
ccaactcnaa ttagatgacc catattcaga aacttactag

40

<210> 37
<211> 54
<212> DNA
<213> Homo sapiens


<400> 37
gtatatatag agttttatat tactttttaa gtacagaatt cataactctca aaaa

54

<210> 38
<211> 41
<212> DNA
<213> Homo sapiens

<400> 38
attgtgacct taattttgtg atctcttgat ttttatttca g

41

 <210> 39
<211> 18
<212> DNA
<213> Homo sapiens

<400> 39
tccccgcctg ccgctctc

18

<210> 40
<211> 18
<212> DNA
<213> Homo sapiens

<400> 40
gcagcggcgg ctcccgtg

18

<210> 41
<211> 20
<212> DNA
<213> Homo sapiens

<400> 41
gtgaacggct ctcagtctgc

20

<210> 42
<211> 19
<212> DNA
<213> Homo sapiens

<400> 42
acgtgcgggg aggaatgga

19

<210> 43
<211> 24
<212> DNA
<213> Homo sapiens

<400> 43
atgatatctt accaaatgat atac

24

<210> 44
<211> 23
<212> DNA
<213> Homo sapiens

<400> 44
ttattcctac ttcttctata cag

23

<210> 45
<211> 21
<212> DNA
<213> Homo sapiens

<400> 45
taccatgct ggctcttttt c

21

c4
<210> 46
<211> 20
<212> DNA
<213> Homo sapiens

<400> 46
tggggccatc ttgttcctga

20

<210> 47
<211> 22
<212> DNA
<213> Homo sapiens

<400> 47
acattaggca caaagcttgc aa

22

<210> 48
<211> 22
<212> DNA
<213> Homo sapiens

<400> 48
atcaagctcc agtaagaagg ta

22

<210> 49
<211> 19
<212> DNA
<213> Homo sapiens

<400> 49
tgcggtcct gggttgttg

19

<210> 50
<211> 20
<212> DNA
<213> Homo sapiens

<400> 50
gccccctcct ttctgaggac

20

<210> 51
<211> 21
<212> DNA
<213> Homo sapiens

<400> 51
ttttctcctg cctcttactg c

21

<210> 52
<211> 20
<212> DNA
<213> Homo sapiens

c4
<400> 52
atgacacccc ccattccctc

20

<210> 53
<211> 24
<212> DNA
<213> Homo sapiens

<400> 53
ccacttaaag cacatatatt tagt

24

<210> 54
<211> 22
<212> DNA
<213> Homo sapiens

<400> 54
gtatggaaaa tagtgaagaa cc

22

<210> 55
<211> 24
<212> DNA
<213> Homo sapiens

<400> 55
ttcttaagtc ctgtttttct ttg

24

<210> 56
<211> 23
<212> DNA

<213> Homo sapiens

<400> 56
tttagaacct tttttgtgtt gtg 23

<210> 57
<211> 24
<212> DNA
<213> Homo sapiens

<400> 57
ctcagattat acactaagcc taac 24

<210> 58
<211> 22
<212> DNA
<213> Homo sapiens

<400> 58
catgtctctt acagtagtac ca 22

<210> 59
<211> 20
<212> DNA
<213> Homo sapiens

<400> 59
aggtccaagg gtagccaagg 20

<210> 60
<211> 27
<212> DNA
<213> Homo sapiens

<400> 60
taaaaatgga taaactacaa ttaaaag 27

<210> 61
<211> 24
<212> DNA
<213> Homo sapiens

<400> 61
aaatacagaa tcatgtcttg aagt 24

<210> 62
<211> 23
<212> DNA
<213> Homo sapiens

<400> 62
acacctaaag atgacaattt gag 23

<210> 63
<211> 24

<212> DNA
 <213> Homo sapiens

 <400> 63 24
 taacttagat agcagtaatt tccc

 <210> 64
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 64 23
 acaataaaact ggagtacaca agg

 <210> 65
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 65 23
 ataggtcatt gcttcttgct gat

 <210> 66
 <211> 24
 <212> DNA
 <213> Homo sapiens

 <400> 66 24
 tgaattttaa tggattacct aggt

 <210> 67
 <211> 25
 <212> DNA
 <213> Homo sapiens

 <400> 67 25
 ctttttttgc ttttactgat taacg

 <210> 68
 <211> 27
 <212> DNA
 <213> Homo sapiens

 <400> 68 27
 tgtaattcat tttattccta atacctc

 <210> 69
 <211> 24
 <212> DNA
 <213> Homo sapiens

 <400> 69 24
 ggtagccata gtatgattat ttct

 <210> 70

<211> 24
<212> DNA
<213> Homo sapiens

<400> 70
ctacctattt ttataccac aaac

24

<210> 71
<211> 23
<212> DNA
<213> Homo sapiens

<400> 71
aagaaagcct acaccatttt tgc

23

<210> 72
<211> 23
<212> DNA
<213> Homo sapiens

<400> 72
gatcattctt agaaccatct tgc

23

<210> 73
<211> 24
<212> DNA
<213> Homo sapiens

<400> 73
acctatagtc taaattatac catc

24

<210> 74
<211> 20
<212> DNA
<213> Homo sapiens

<400> 74
gtcatggcat tagtgaccag

20

<210> 75
<211> 24
<212> DNA
<213> Homo sapiens

<400> 75
agtcgtaatt ttgtttctaa actc

24

<210> 76
<211> 21
<212> DNA
<213> Homo sapiens

<400> 76
tgaaggactc ggatttcacc c

21

<210> 77
<211> 23
<212> DNA
<213> Homo sapiens

<400> 77
tcattcactc acagcctgat gac

23

<210> 78
<211> 22
<212> DNA
<213> Homo sapiens

<400> 78
gctttgaaac atgcactacg at

22

<210> 79
<211> 24
<212> DNA
<213> Homo sapiens

<400> 79
aaacatcatt gctcttcaaa taac

24

c64
<210> 80
<211> 24
<212> DNA
<213> Homo sapiens

<400> 80
taccatgatt taaaaatcca ccag

24

<210> 81
<211> 23
<212> DNA
<213> Homo sapiens

<400> 81
gatgattgtc tttttcctct tgc

23

<210> 82
<211> 24
<212> DNA
<213> Homo sapiens

<400> 82
ctgagctatc ttaagaaata catg

24

<210> 83
<211> 25
<212> DNA
<213> Homo sapiens

<400> 83
ttttaaatga tcctctattc tgtat

25

<210> 84
<211> 24
<212> DNA
<213> Homo sapiens

<400> 84
acagagtcag accctgcctc aaag

24

<210> 85
<211> 23
<212> DNA
<213> Homo sapiens

<400> 85
tttctattct tactgctagc att

23

<210> 86
<211> 22
<212> DNA
<213> Homo sapiens

<400> 86
atacacaggt aagaaattag ga

22

cf
<210> 87
<211> 22
<212> DNA
<213> Homo sapiens

<400> 87
tagatgaccc atattctggt tc

22

<210> 88
<211> 22
<212> DNA
<213> Homo sapiens

<400> 88
caattaggtc tttttgagag ta

22

<210> 89
<211> 22
<212> DNA
<213> Homo sapiens

<400> 89
gttactgcat acacattgtg ac

22

<210> 90
<211> 23
<212> DNA
<213> Homo sapiens

<400> 90

gctttttgtt tcctaacatg aag

23

<210> 91
<211> 21
<212> DNA
<213> Homo sapiens

<400> 91
tctcccacag gtaatactcc c

21

<210> 92
<211> 21
<212> DNA
<213> Homo sapiens

<400> 92
gctagaactg aatggggtac g

21

<210> 93
<211> 22
<212> DNA
<213> Homo sapiens

<400> 93
caggacaaaa taatcctgtc cc

22

<210> 94
<211> 24
<212> DNA
<213> Homo sapiens

<400> 94
attttcttag tttcattctt cctc

24

<210> 95
<211> 25
<212> DNA
<213> Homo sapiens

<400> 95
agaaggatcc cttgtgcagt gtgga

25

<210> 96
<211> 24
<212> DNA
<213> Homo sapiens

<400> 96
gacaggatcc tgaagctgag ttg

24

<210> 97
<211> 18
<212> DNA
<213> Homo sapiens

<400> 97 18
 tcagaaagtg ctgaagag

 <210> 98
 <211> 19
 <212> DNA
 <213> Homo sapiens

 <400> 98 19
 ggaataatta ggtctccaa

 <210> 99
 <211> 21
 <212> DNA
 <213> Homo sapiens

 <400> 99 21
 gcaaataccta agagagaaca a

 <210> 100
 <211> 19
 <212> DNA
 <213> Homo sapiens

 <400> 100 19
 gatggcaagc ttgagccag

 <210> 101
 <211> 18
 <212> DNA
 <213> Homo sapiens

 <400> 101 18
 gttccagcag tgtcacag

 <210> 102
 <211> 18
 <212> DNA
 <213> Homo sapiens

 <400> 102 18
 gggagatttc gctcctga

 <210> 103
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 103 23
 agtacaagga tgccaatatt atg

 <210> 104
 <211> 23
 <212> DNA
 <213> Homo sapiens

<400> 104 23
 acttctatct ttttcagaac gag

 <210> 105
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 105 23
 atttgaatac tacagtgtta ccc

 <210> 106
 <211> 24
 <212> DNA
 <213> Homo sapiens

 <400> 106 24
 cttgtattct aatttggcat aagg

 <210> 107
 <211> 22
 <212> DNA
 <213> Homo sapiens

 CF <400> 107 22
 ctgcccatac acattcaaac ac

 <210> 108
 <211> 21
 <212> DNA
 <213> Homo sapiens

 <400> 108 21
 tgtttgcgtc ttgcccattct t

 <210> 109
 <211> 24
 <212> DNA
 <213> Homo sapiens

 <400> 109 24
 agtctttaaatt attcagatga gcag

 <210> 110
 <211> 26
 <212> DNA
 <213> Homo sapiens

 <400> 110 26
 gtttctcttc atttatatttt atgcta

 <210> 111
 <211> 23
 <212> DNA

<213> Homo sapiens

<400> 111
aagcctacca attatagtga acg

23

<210> 112
<211> 23
<212> DNA
<213> Homo sapiens

<400> 112
agctgatgac aaagatgata atc

23

<210> 113
<211> 24
<212> DNA
<213> Homo sapiens

<400> 113
aagaaacaat acagacttat tgtg

24

<210> 114
<211> 20
<212> DNA
<213> Homo sapiens

<400> 114
atgagtgggg tctcctgaac

20

<210> 115
<211> 21
<212> DNA
<213> Homo sapiens

<400> 115
atctccctcc aaaagtgggtg c

21

<210> 116
<211> 22
<212> DNA
<213> Homo sapiens

<400> 116
tccatctgga gtactttctg tg

22

<210> 117
<211> 22
<212> DNA
<213> Homo sapiens

<400> 117
agtaaagtct gcagttcaga gg

22

<210> 118
<211> 19

<212> DNA
 <213> Homo sapiens

 <400> 118 19
 ccgtggcata tcatcccc

 <210> 119
 <211> 22
 <212> DNA
 <213> Homo sapiens

 <400> 119 22
 cccagactgc ttcaaaatta cc

 <210> 120
 <211> 21
 <212> DNA
 <213> Homo sapiens

 <400> 120 21
 gagcctcatc tgtacttctg c

 <210> 121
 <211> 21
 <212> DNA
 <213> Homo sapiens

 <400> 121 21
 ccctccaaat gagttagctg c

 <210> 122
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 122 23
 ttgtggtata ggttttactg gtg

 <210> 123
 <211> 23
 <212> DNA
 <213> Homo sapiens

 <400> 123 23
 acccaacaaa aatcagttag atg

 <210> 124
 <211> 21
 <212> DNA
 <213> Homo sapiens

 <400> 124 21
 gtggctggta actttagcct c

 <210> 125

<211> 21
<212> DNA
<213> Homo sapiens

<400> 125
atgatgttga cttttccagg g

21

<210> 126
<211> 24
<212> DNA
<213> Homo sapiens

<400> 126
attgtgtaac ttttcatcag ttgc

24

<210> 127
<211> 21
<212> DNA
<213> Homo sapiens

<400> 127
aaagacatac cagacagagg g

21

<210> 128
<211> 21
<212> DNA
<213> Homo sapiens

<400> 128
cttttttggc attgcggagc t

21

<210> 129
<211> 22
<212> DNA
<213> Homo sapiens

<400> 129
aagatgacct gttgcaggaa tg

22

<210> 130
<211> 24
<212> DNA
<213> Homo sapiens

<400> 130
gaatcagacc aagcttgtct agat

24

<210> 131
<211> 24
<212> DNA
<213> Homo sapiens

<400> 131
caatagtaag tagtttacat caag

24

<210> 132
<211> 22
<212> DNA
<213> Homo sapiens

<400> 132
aaacaggact tgtactgtag ga

22

<210> 133
<211> 21
<212> DNA
<213> Homo sapiens

<400> 133
cagccccttc aagcaaaca c

21

<210> 134
<211> 22
<212> DNA
<213> Homo sapiens

<400> 134
gaggacttat tccatttcta cc

22

<210> 135
<211> 20
<212> DNA
<213> Homo sapiens

<400> 135
cagtctcctg gccgaaactc

20

<210> 136
<211> 22
<212> DNA
<213> Homo sapiens

<400> 136
gttgactggc gtactaatac ag

22

<210> 137
<211> 23
<212> DNA
<213> Homo sapiens

<400> 137
tggtaatgga gccataaaa agg

23

<210> 138
<211> 20
<212> DNA
<213> Homo sapiens

<400> 138
tgggactttt cgccatccac

20

<210> 139
<211> 22
<212> DNA
<213> Homo sapiens

<400> 139 22
tgtctctatc cacacattcg tc

<210> 140
<211> 24
<212> DNA
<213> Homo sapiens

<400> 140 24
atgtttttca tcctcacttt ttgc

<210> 141
<211> 22
<212> DNA
<213> Homo sapiens

<400> 141 22
ggagaagaac tggaagttca tc

cf
<210> 142
<211> 25
<212> DNA
<213> Homo sapiens

<400> 142 25
ttgaatcttt aatgtttgga tttgc

<210> 143
<211> 21
<212> DNA
<213> Homo sapiens

<400> 143 21
tctccacag gtaatactcc c

<210> 144
<211> 21
<212> DNA
<213> Homo sapiens

<400> 144 21
gctacaactg aatgggggtac g

<210> 145
<211> 22
<212> DNA
<213> Homo sapiens

<400> 145

caggacaaaa taatcctgtc cc

22

<210> 146
<211> 24
<212> DNA
<213> Homo sapiens

<400> 146
attttctttac tttcattctt cctc

24

<210> 147
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> consensus sequence

<400> 147
Phe Val Glu Thr Pro Cys Phe Ser Arg Ser Ser Leu Ser Ser Leu Ser
1 5 10 15

<210> 148
<211> 20
<212> PRT
<213> Homo sapiens

<400> 148
Tyr Cys Val Glu Asp Thr Pro Ile Cys Phe Ser Arg Cys Ser Ser Leu
1 5 10 15
Ser Ser Leu Ser
20

<210> 149
<211> 20
<212> PRT
<213> Homo sapiens

<400> 149
His Thr Val Gln Glu Thr Pro Leu Met Phe Ser Arg Cys Thr Ser Val
1 5 10 15
Ser Ser Leu Asp
20

<210> 150
<211> 20
<212> PRT
<213> Homo sapiens

<400> 150
Phe Ala Thr Glu Ser Thr Pro Asp Gly Phe Ser Cys Ser Ser Ser Leu
1 5 10 15
Ser Ala Leu Ser
20

<210> 151
<211> 20
<212> PRT
<213> Homo sapiens

<400> 151
Tyr Cys Val Glu Gly Thr Pro Ile Asn Phe Ser Thr Ala Thr Ser Leu
1 5 10 15
Ser Asp Leu Thr
20

<210> 152
<211> 20
<212> PRT
<213> Homo sapiens

<400> 152
Thr Pro Ile Glu Gly Thr Pro Tyr Cys Phe Ser Arg Asn Asp Ser Leu
1 5 10 15
Ser Ser Leu Asp
20

<210> 153
<211> 20
<212> PRT
<213> Homo sapiens

104
<400> 153
Phe Ala Ile Glu Asn Thr Pro Val Cys Pro Ser His Asn Ser Ser Leu
1 5 10 15
Ser Ser Leu Ser
20

<210> 154
<211> 20
<212> PRT
<213> Homo sapiens

<400> 154
Arg His Val Glu Asp Thr Pro Val Cys Phe Ser Arg Asn Ser Ser Leu
1 5 10 15
Ser Ser Leu Ser
20